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***RET/PTC* and *PAX8/PPAR γ* chromosomal rearrangements in post-Chernobyl thyroid cancer and their association with I-131 radiation dose and other characteristics**

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Abstract

Background—Childhood exposure to I-131 from the 1986 Chernobyl accident led to a sharp increase in papillary thyroid carcinoma (PTC) incidence in regions surrounding the reactor. Data concerning the association between genetic mutations in PTCs and individual radiation doses are limited.

Methods—We performed mutational analysis of 62 PTCs diagnosed in a Ukrainian cohort of patients who were <18 y.o. in 1986 and received 0.008-8.6 Gy of I-131 to the thyroid and explored associations between mutation types and I-131 dose and other characteristics.

Results—*RET/PTC* rearrangements were most common (35%), followed by *BRAF* (15%) and *RAS* (8%) point mutations. Two tumors carrying *PAX8/PPAR γ* rearrangement were identified. We found a significant negative association with I-131 dose for *BRAF* and *RAS* point mutations and a significant concave association with I-131 dose, with an inflection point at 1.6 Gy and odds ratio 2.1, based on a linear-quadratic model for *RET/PTC* and *PAX8/PPAR γ* rearrangements. The trends with dose were significantly different between tumors with point mutations and rearrangements. Compared to point mutations, rearrangements were associated with residence in the relatively iodine deficient Zhytomyr region, younger age at exposure or surgery, and male gender.

Conclusions—Our results provide the first demonstration of *PAX8/PPAR γ* rearrangements in post-Chernobyl tumors and show different associations for point mutations and chromosomal rearrangements with I-131 dose and other factors. These data support the relationship between chromosomal rearrangements, but not point mutations, and I-131 exposure and point to a possible role of iodine deficiency in generation of *RET/PTC* rearrangements in these patients.

Keywords

Chernobyl; papillary thyroid carcinoma; iodine-131; *RET/PTC*; *PAX8/PPAR γ*

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INTRODUCTION

Exposure to ionizing radiation during childhood is known to cause thyroid cancer, with significant dose-dependently increased incidence, particularly in children and young adults.¹ After the Chernobyl accident of 1986, residents of regions surrounding the Chernobyl nuclear power plant, including Ukraine, Belarus, and the Russian Federation, received variable doses of radioiodines through inhalation and ingestion of contaminated dairy products or vegetables. These regions experienced a dramatic rise in incidence of thyroid cancer,² with at least 5,000 new cases observed in individuals exposed during childhood or adolescence.³ Papillary thyroid carcinoma (PTC) is known to be the principal type of thyroid carcinoma associated with radiation exposure and comprised the majority of pediatric thyroid tumors in residents of the regions surrounding Chernobyl.^{2,4,5}

Case-control and cohort studies of post-Chernobyl thyroid cancers have demonstrated that the risk of thyroid carcinoma is strongly related to I-131 dose absorbed by the thyroid.⁶⁻⁹ The reported excess relative risk per unit of dose (Gy) are between 2 to 5. Additionally, age at exposure and iodine deficiency have been found to modify the I-131 related risk of thyroid cancer with higher risk per unit of dose observed in persons exposed as younger children, particularly infants,¹⁰ and in individuals living in areas with low soil iodine content.⁷

Somatic genetic alterations that activate the mitogen activated protein kinase (MAPK) signaling cascade are known to be “driver” mutations that play a crucial role in the development of PTC.¹¹ These include point mutations in *BRAF* and *RAS* as well as chromosomal rearrangements such as *RET/PTC*. Whereas MAPK activation via point mutations is far more common in sporadic thyroid cancer, in tumors associated with radiation exposure, this pathway is most frequently activated via *RET/PTC* and other chromosomal rearrangement. Studies of post-Chernobyl and post-radiotherapy tumors have found *RET/PTC* rearrangements in up to of PTCs.¹²⁻¹⁴ The link between chromosomal rearrangements and exposure to ionizing radiation has also been supported by studies that have demonstrated induction of *RET/PTC* in human thyroid cell lines and tissue xenografts in SCID mice by X-ray or γ -radiation.^{15,16} Recent studies have led to better understanding of mechanisms by which radiation exposure induces chromosomal rearrangements. Studies of both *RET/PTC* and *TRK* rearrangements have shown that the gene loci involved in fusions lie in close spatial proximity to one another within the human thyroid cell nucleus at the time of exposure,¹⁷⁻¹⁹ likely predisposing to recombination of adjacent chromosomal regions radiation-induced DNA damage.

However, the association of chromosomal rearrangements or other mutational events with individual radiation doses in humans is not well established. Among PTCs that individuals exposed to predominantly γ -radiation from the atomic bombings in Hiroshima and Nagasaki, higher doses were associated with higher prevalence of *RET/PTC* rearrangements and lower prevalence of *BRAF* point mutations.^{20,21} By contrast, no significant association between *RET/PTC* activation and individual I-131 doses was found in one post-Chernobyl study of cancer occurring in the Bryansk oblast of the Russian Federation.²² The prevalence of another rearrangement type, *PAX8/PPAR γ* , known to occur in follicular thyroid cancer and only occasionally found in PTC,¹¹ has not been studied in post-Chernobyl tumors.

Herein, we report the results of mutational analysis of a series of post-Chernobyl PTCs from Ukrainian individuals with measurement-based estimates of I-131 dose to the thyroid reconstructed as part of the Ukrainian-American cohort study.²³ Based on the aforementioned experimental and human data, we hypothesized that *RET/PTC*

rearrangements represent a common genetic events in these cancers and chromosomal rearrangements and point mutations have different association with I-131 dose. The obtained results demonstrate the predominance of chromosomal rearrangements in these tumors, show for the first time the occurrence of *PAX8/PPAR γ* rearrangements in post-Chernobyl tumors, and establish associations of specific genetic alterations with I-131 doses and other patient characteristics.

MATERIALS AND METHODS

Patients and tissue samples

Cases included patients who were part of the Ukrainian-American cohort study and underwent surgery for suspected thyroid carcinoma.²³ The cohort is composed of 13,243 Ukrainian residents, less than 18 years old at the time of the Chernobyl accident, with individual radioactivity measurements taken within two months after the accident. After four sequential screening examinations, 110 thyroid carcinomas, including 104 PTCs, were diagnosed between 1998 and 2008 at the Laboratory of Morphology of Endocrine System of the Institute of Endocrinology and Metabolism (IEM, Kyiv, Ukraine).²⁴ The International Pathology Panel, within the Chernobyl Tissue Bank (CTB), reviewed all pathological diagnoses. Seventy five of 104 cases of PTC had at least one frozen tissue specimen from which DNA or RNA were extracted at IEM or Imperial College (London, UK). Nucleic acids from 74 PTCs were received through the CTB. Four cases from a separate cohort exposed *in utero* were excluded. Eight cases that lacked either DNA (n=3) or RNA (n=5) were also excluded.

Estimation of I-131 thyroid doses

Dosimetric methods have been described in detail.^{25,26} Briefly, individual I-131 thyroid doses and their uncertainties were estimated from thyroid radioactivity measurements, data on dietary and lifestyle habits, and environmental transfer models using a Monte-Carlo procedure with 1,000 realizations per individual.²⁶ For the analysis, we used the arithmetic mean of each individual's 1,000 realizations as the best estimate of I-131 dose corrected for thyroid masses typical of the Ukrainian population.⁶

Nucleic Acids

Tumor DNA and RNA were obtained through CTB. The samples were received with a CTB code that was later linked with a code for the individual in the Ukrainian-American study.²³

Detection of point mutations

All tumors for which DNA was available were tested for point mutations in *BRAF* (V600E and K601E), *NRAS* (codon 61), *HRAS* (codon 61), and *KRAS* (codons 12/13) genes using fluorescence melting curve analysis as previously described.²⁷ Briefly, the samples were amplified on the LightCycler (Roche Diagnostics, Indianapolis, IN) using the LightCycler FASTstart DNA Master Mix (Roche) and specific probes.²⁷ Post-PCR melting curves were compared to those from control tumors known to have/lack specific point mutations. All mutations were confirmed by Sanger sequencing.

Detection of chromosomal rearrangements

All tumors for which RNA was available were screened for *RET/PTC1*, *RET/PTC3*, and *PAX8/PPAR γ* using real-time RT-PCR. Tumor RNA was reverse transcribed and amplified on the 7500 Real Time PCR System (Applied Biosystems) using the Taqman Universal PCR Master Mix (Applied Biosystems) and specific probes.²⁷ For *PAX8/PPAR γ* , a Δ Ct of <10 cycles, as compared to the amplification of *GAPDH*, was used as a cut-off. For *PAX8/*

PPAR γ and *RET/PTC3*, the presence of the rearrangement was confirmed by conventional RT-PCR and agarose gel electrophoresis. All chromosomal rearrangements were confirmed by Sanger sequencing.

Statistical analysis

Mutation prevalence data was analyzed using standard logistic regression modeling. The probability of a *BRAF* or *RAS* positive mutation for the relevant endpoint given effect modifying variables: age at surgery *a*, gender *s*, oblast *O* and Chernobyl-associated radiation dose *D* (in Gy), is given by:

$$P[\text{mutation}|D, a, s] = \frac{\exp[\alpha_0 + \alpha_1(a - 25) + \alpha_2 1_{\text{gender=female}} + \alpha_3 1_{O=\text{zhytomir}} + \alpha_4 1_{O=\text{Kyiv}} + \alpha_5 D]}{1 + \exp[\alpha_0 + \alpha_1(a - 25) + \alpha_2 1_{\text{gender=female}} + \alpha_3 1_{O=\text{zhytomir}} + \alpha_4 1_{O=\text{Kyiv}} + \alpha_5 D]}$$

The model is fitted by binomial maximum likelihood²⁸ using Epicure. Age (at surgery) was centered by subtracting 25 years to aid convergence of fitted models. For chromosomal rearrangements, the linear and quadratic terms for dose (dose+dose²) were used, while for point mutations a linear model in dose sufficed. Unless otherwise stated, all confidence intervals were profile-likelihood based.²⁸ All tests were two-sided and based on the likelihood-ratio test.²⁹ Adjustments were made for age at surgery, gender, oblast of residence at the time of screening and dose because of indications of significant or borderline significant effects on mutation rates. Oblast at the time of screening compared with that at the time of exposure differed for only three patients. Tests of heterogeneity were performed as described by Pierce and Preston.³⁰ Analysis of time from exposure to surgery used a standard linear regression model, using log-transformed time since exposure. All linear regression analyses were performed using Stata. Mean age at exposure was compared by two sample t-test.

RESULTS

Case characteristics

The case series includes 26 males (42%) and 36 females (58%) living in areas surrounding Chernobyl, i.e. the Zhytomyr (n=17; 27.4%), Kyiv (n=12; 19.4%) or Chernihiv (n=33; 53.2%) oblasts, who were between 5 months and 17 years old (mean, 8.0 years) at the time of the Chernobyl accident. The estimated I-131 dose for patients in the study ranged from 0.008 Gy to 8.6 Gy, with a mean dose of 1.3 Gy. Surgical removal of PTCs occurred between October, 1998 and December, 2007, with time between exposure and surgery ranging from 12.5 to 21.6 years (mean, 16.5 years).

Mutation analysis

RET/PTC rearrangement was the most common genetic alteration and was found in 22 (35%) cases, including 14 *RET/PTC1* and 8 *RET/PTC3* rearrangements (Table 1). Point mutations in the *BRAF* and *RAS* genes were found in 9 (15%) and 5 (8%) of the tumors, respectively. All nine *BRAF* mutations were V600E. Four *RAS* mutations were detected in *NRAS* codon 61 and one in *HRAS* codon 61. No *KRAS* mutations were found in codons 12 and 13. Additionally, we studied these tumors for *PAX8/PPAR γ* rearrangement, a prototypic genetic alteration found in thyroid follicular carcinoma that occurs with lower prevalence in the follicular variant of PTC. Two tumors were positive for *PAX8/PPAR γ* , both were of the follicular variant of PTC. In both cases, the fusions were between exon 9 of *PAX8* and exon 1 of *PPAR γ* , with several expected splice variants of the chimeric *PAX8/PPAR γ* transcripts detected. One tumor had more than one mutation, harboring an *NRAS*

point mutation in codon 61 and *PAX8/PPAR γ* rearrangement. Twenty-five (40%) tumors revealed none of the studied mutations.

Univariate analysis of mutation type and exposure-related characteristics

Patients with tumors positive for *BRAF* or *RAS* point mutations had the lowest average dose of I-131 (0.27 Gy and 0.20 Gy, respectively), significantly lower than that for all other patients ($p < 0.001$). Patients with tumors harboring *RET/PTC1* or *RET/PTC3* rearrangements received average doses of 1.04 Gy and 1.54 Gy, respectively. Patients with tumors negative for any of these mutations had the highest average dose (1.97 Gy). Additionally, as compared to all other patients, patients with tumors harboring *RET/PTC1* or *RET/PTC3* were significantly younger at the time of exposure (6.4 and 6.3 years, respectively) whereas patients with *BRAF* or *RAS* mutations were significantly older at the time of exposure (10.2 and 10.9 years, respectively) ($p = 0.01$ for both). In our case series, age at exposure correlated with age at surgery (or attained age) (Pearson correlation coefficient, $r^2 = 0.85$). Thus, patients with tumors having *RET/PTC1* or *RET/PTC3* rearrangements were also younger at the time of surgery (mean age 23.6 years and 20.1 years, respectively) ($p = 0.007$) and patients with tumors positive for *BRAF* or *RAS* mutations were older (mean age 27.1 years and 29.4 years, respectively) ($p = 0.002$) than other cases. The mean time between exposure and surgery for *RET/PTC3* positive cases was 13.7 years, significantly shorter than that for all other cases combined ($p < 0.001$) (Table 1).

Multivariate analysis of tumors positive for *BRAF* or *RAS* point mutations

Factors independently associated with tumors harboring *BRAF* or *RAS* point mutations, as compared to all other tumors, are shown in Table 2. Adjusting for age at surgery, gender, and oblast of residence, there was a significant negative association between these point mutations and I-131 dose ($p = 0.001$). The estimated regression coefficient for the dose term based on a log-linear model was -2.51 per Gy (95% CI: $-5.42, -0.78$) (Fig. 1A). In addition, the point mutations were associated with older age at surgery ($p = 0.014$) and female gender ($p = 0.002$).

Multivariate analysis of tumors positive for *RET/PTC* or *PAX8/PPAR γ* chromosomal rearrangements

There was a significant negative dose-response relationship using continuous I-131 dose ($p = 0.040$, not shown), which was improved at marginal levels of statistical significance ($p = 0.053$) by addition of a quadratic term in dose (dose²); overall the linear-quadratic dose response was statistically significant (2 df trend, $p = 0.019$; Table 3) with estimated regression coefficients of 0.96 per Gy (95% CI: $-0.54, 2.61$) for the linear term and of -0.31 per Gy² for the quadratic term (95% CI: $-0.70, 0.00$). While the parameters of the dose-response function are imprecise, as evidenced by wide confidence intervals, the data suggest a non-monotone relationship for *RET/PTC* or *PAX8/PPAR γ* -positive tumors and dose with increased risk at low-to-moderate doses and decreased risk at high doses (Fig. 1B). The latter may reflect the fact that cases with no known mutation were associated with the highest average dose overall ($p = 0.012$). Indeed, among the patients who received doses above 1.6 Gy, 10 out of 16 (62.5%), had tumors that were negative for all mutations studied.

In addition to dose, the presence of chromosomal rearrangement was positively associated with residence in the Zhytomyr oblast ($p = 0.007$ for 2 df test of oblast differences) and negatively associated with female gender ($p = 0.043$) and attained age ($p = 0.001$) (Table 3). The presence of *RET/PTC3* rearrangement compared to all other tumors was negatively associated with time between exposure and surgery, even when adjusted for I-131 dose and age at exposure ($p = 0.001$) or attained age ($p = 0.012$, not shown). No association with time from exposure to surgery was found for any other rearrangements or point mutations.

Comparison between tumors positive for point mutations and those harboring chromosomal rearrangements

Direct comparison of tumors with point mutations (*BRAF* or *RAS*) and chromosomal rearrangements (*RET/PTC* or *PAX8/PPAR γ*) using multivariate logistic regression demonstrated a significant difference in trends with I-131 dose ($p=0.020$; Table 4). Additionally, age at surgery ($p<0.001$), gender ($p<0.001$), and oblast of residence ($p=0.003$) were significantly and independently associated with presence of point mutations relative to chromosomal rearrangements. Compared to *BRAF* or *RAS* positive tumors, *RET/PTC* or *PAX8/PPAR γ* positive tumors were likely to occur in individuals with higher I-131 doses, younger age at surgery (and therefore younger at exposure), in males, and residents of Zhytomyr oblast (Table 4).

DISCUSSION

Our study of post-Chernobyl thyroid tumors confirm the high frequency of chromosomal rearrangements, particularly *RET/PTC*, and lower frequency of *BRAF* and *RAS* point mutations, compared to those typically observed in sporadic tumors,¹¹ and report for the first time the occurrence of *PAX8/PPAR γ* rearrangements in post-Chernobyl PTCs. Moreover, we identified significant independent differences between chromosomal rearrangements and point mutations with respect to I-131 thyroid doses, gender, oblast of residence, and age at exposure or surgery, suggesting that these tumors have distinct etiologies, i.e. tumors with chromosomal rearrangements, but not with point mutations, are likely to be radiation-related.

Unique features of this study include individual I-131 thyroid doses based on radioactivity measurements,^{6,25} well-characterized tumors detected through standardized medical examinations, and comprehensive molecular profiling. Consequently, we were able to draw associations between specific mutation types and I-131 dose and other contributing factors. The association between I-131 thyroid dose and presence of chromosomal rearrangements followed a linear quadratic function, indicating a positive relationship in the low-to-moderate dose range and a negative relationship at high doses. It is likely that tumors with no known mutations accounted for the observed downturn at high doses, as their proportion relative to other tumors reached 62% at doses of 1.6 Gy or higher. Interestingly, a positive linear dose-response for *RET/PTC* rearrangements has been reported in thyroid tumors that developed among the atomic bomb survivors of Hiroshima and Nagasaki,^{20,21} but very few cases in these studies received doses higher than 1.6 Gy.²⁰ No significant association was found for *RET/PTC* activation with individual I-131 doses in the study of PTCs from Bryansk oblast of the Russian Federation.²² However, these patients were exposed to lower doses (mean 0.363 Gy for childhood cancers and 0.039 Gy for adult cancers) than those in the current study (mean 1.3 Gy). The inconsistencies in dose-response findings across these studies are likely to result from different dose and age range, geographic origin, uncertainties in dose estimates, or limited sample size.

By contrast, patients with tumors harboring *BRAF* and *RAS* point mutations received the lowest average I-131 doses, were oldest at the time of the Chernobyl accident or at surgery, and were predominantly female. The strong negative association for point mutation-positive tumors with dose found when comparing *BRAF* and *RAS* positive tumors against all others is consistent with that in atomic bomb survivors.²⁰ The negative association with dose together with the fact that the *BRAF* mutation is most commonly found in sporadic thyroid cancer,^{11,31} incidence of which rapidly increases during the third decade of life and is three times as common in women,^{32,33} suggest that *BRAF* and *RAS* positive tumors found in this cohort were likely to develop via pathogenic mechanisms more typical of sporadic thyroid cancer.

More than one-third of the tumors in our study had no identifiable mutations, and exhibited their own unique characteristics. The age at exposure or surgery for individuals with such tumors was higher than that of patients with *RET/PTC* rearrangements, but lower than that of patients with point mutations. These patients also received the highest I-131 thyroid doses. Therefore, the development of these tumors is likely to be related to radiation exposure, but involved other, unknown mutations. A recent study of PTCs in atomic bomb survivors found *ALK* rearrangements, although they were detected at very low levels and using highly sensitive analyses, leaving the biological significance of this finding unclear.³⁴ Other chromosomal rearrangements that occur very rarely in sporadic PTC but have been seen with higher frequency in PTCs following radiation exposure include *BRAF/AKAP9* and *TRK* rearrangements.¹¹ It is possible that these unmeasured, rare rearrangements may partially compose the set of tumors with as yet unknown mutations in our study.

Another unexpected finding in the current study was a strong association between *RET/PTC* and residence in Zhytomyr oblast. Although study participants from Zhytomyr received higher I-131 doses than those of the neighboring Ukrainian oblasts, the association with *RET/PTC* was independent of dose. The Zhytomyr oblast has no noticeable geographic or ethnic differences, but is associated with relative iodine deficiency. Indeed, residents of Zhytomyr have been shown to have lower levels of urinary iodine excretion than those of Kyiv or Chernihiv oblasts.³⁵ Iodine deficiency has been shown to contribute to the risk of post-Chernobyl thyroid cancer in Belarus,^{3,7} and may possibly be responsible for the increased frequency of *RET/PTC* in tumors from Zhytomyr oblast residents. It is conceivable that more avid trapping and intracellular metabolism of I-131 by thyroid cells under conditions of high thyroid stimulating hormone (TSH) stimulation produce more extensive damage to the nuclear DNA, resulting in *RET/PTC* rearrangement. If confirmed, this would provide a biological basis for the higher risk of radiation-induced thyroid cancer in areas of relative iodine deficiency.

This study also identified *PAX8/PPAR γ* rearrangements in post-Chernobyl tumors. This rearrangement is known to occur with high frequency in another type of thyroid cancer, follicular carcinoma, and with much lower frequency in the follicular variant of PTC.¹¹ Both tumors positive for *PAX8/PPAR γ* rearrangement in our study were follicular variant of PTC. In one large study, tumors with *PAX8/PPAR γ* rearrangements were more common in patients with a history of preceding non-thyroid cancer,³⁶ which may implicate therapeutic radiation. However, to our knowledge, *PAX8/PPAR γ* has not been previously reported in post-Chernobyl or other radiation-associated cancers.

In summary, this study provides strong support for the association between chromosomal rearrangements and exposure to I-131 from the Chernobyl accident. Furthermore, our findings point to a possible role of iodine deficiency in generation of *RET/PTC* rearrangements. Finally, since a significant proportion of tumors in our study had no detectable mutations and were associated with high radiation doses, we hypothesize that undiscovered mutational events important in radiation-induced thyroid carcinogenesis must exist.

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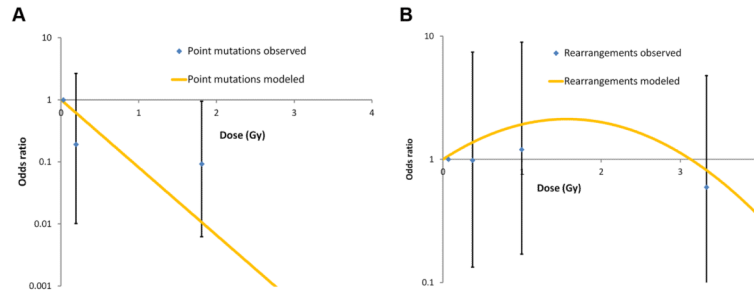


Figure 1. Dose-response relationship for point mutations and chromosomal rearrangements

A. Dose-response relationship for patients with tumors harboring *BRAF* or *RAS* point mutations. Odds ratio data for the relationship between chromosomal rearrangement and estimated I-131 dose fitted to a linear quadratic model with adjustments for a sex, oblast, age of surgery, and dose. **B.** Dose-response relationship for patients with tumors harboring *RET/PTC* or *PAX8/PPAR γ* chromosomal rearrangements. Odds ratio data for the relationship between chromosomal rearrangement and estimated I-131 dose fitted to a linear quadratic model with adjustments for a sex, oblast, age of surgery, and dose.

Genetic alterations and exposure-related characteristics among cases of thyroid cancer developed after the Chernobyl accident in Ukraine

Table 1

| Genetic Alteration | Mutation frequency | I-131 dose, mean (Gy) | Age at exposure, mean (yr) | Age at surgery, mean (yr) | Time since exposure, mean (yr) |
|---------------------|--------------------|-----------------------|----------------------------|---------------------------|--------------------------------|
| <i>RET/PTC1</i> | 14 (22%) | 1.04 | 6.4 | 23.6 | 16.9 |
| <i>RET/PTC3</i> | 8 (13%) | 1.54 | 6.3 | 20.1 | 13.7 |
| <i>BRAF</i> | 9 (15%) | 0.27 | 10.2 | 27.1 | 16.8 |
| <i>RAS</i> * | 5 (8%) | 0.20 | 10.9 | 29.4 | 18.6 |
| <i>PAX8/PPARY</i> * | 2 (3%) | 0.62 | 12.2 | 25.8 | 13.5 |
| No known mutation | 25 (40%) | 1.97 | 7.8 | 24.4 | 16.6 |
| Total/overall | 62 (100%) | 1.27 | 8.1 | 24.6 | 16.5 |

* One case was found to be positive for both *NRAS* mutation and *PAX8/PPARY* rearrangement.

Table 2Factors associated with *BRAF* or *RAS* point mutations in multivariate analysis

| | <i>BRAF</i> or <i>RAS</i> mutation positive | <i>BRAF</i> or <i>RAS</i> mutation negative | | |
|----------------------------|---|---|--------------------|----------------|
| Characteristic | % or mean (SD) | % or mean (SD) | OR | 95% CI |
| I-131 dose, Gy | | | | |
| 0.008-<0.05 | 35.7 | 3.9 | 1.00 | Referent |
| 0.05-<0.35 | 28.6 | 19.6 | 0.19 | (0.01, 2.67) |
| 0.35-8.60 | 35.7 | 76.5 | 0.09 | (0.01, 0.95) |
| P trend | | | 0.001 [/] | |
| Oblast of residence | | | | |
| Chernihiv | 71.4 | 47.1 | 1.00 | Referent |
| Kyiv | 21.4 | 21.6 | 0.24 | (0.02, 2.33) |
| Zhytomyr | 7.1 | 31.4 | 0.07 | (0.00, 0.82) |
| P heterogeneity | | | 0.086 | |
| Age at surgery, yr | 28.1 (4.2) | 23.4 (4.9) | 1.26 | (1.05, 1.61) |
| P trend | | | 0.014 | |
| Sex | | | | |
| Male | 14.3 | 47.1 | 1.00 | Referent |
| Female | 85.7 | 52.9 | 21.59 | (2.69, 349.32) |
| P | | | 0.002 | |

[/]Based on linear dose-response model.

Table 3Factors associated with *RET/PTC* or *PAX8/PPAR γ* rearrangements in multivariate analysis

| | <i>RET/PTC</i> or <i>PAX8/PPARγ</i> positive | <i>RET/PTC</i> or <i>PAX8/PPARγ</i> negative | | |
|----------------------------|---|---|--------------------|---------------|
| Characteristic | % or mean (SD) | % or mean (SD) | OR | 95% CI |
| I-131 dose, Gy | | | | |
| 0.008-<0.4 | 25.0 | 39.5 | 1.00 | Referent |
| 0.4-<0.91 | 25.0 | 23.7 | 1.38 | (0.28, 7.00) |
| 0.91-<1.63 | 25.0 | 10.5 | 2.47 | (0.34, 19.05) |
| 1.63-8.60 | 25.0 | 26.3 | 0.77 | (0.13, 4.13) |
| P trend | | | 0.019 ¹ | |
| Oblast of residence | | | | |
| Chernihiv | 45.8 | 57.9 | 1.00 | Referent |
| Kyiv | 12.5 | 23.7 | 0.88 | (0.13, 5.13) |
| Zhytomyr | 41.7 | 18.4 | 11.66 | (2.22, 82.52) |
| P heterogeneity | | | 0.007 | |
| Age at surgery, yr | 22.6 (4.5) | 25.8 (5.3) | 0.79 | (0.66, 0.91) |
| P trend | | | 0.001 | |
| Sex | | | | |
| Male | 54.2 | 36.8 | 1.00 | Referent |
| Female | 45.8 | 63.2 | 0.27 | (0.06, 0.96) |
| P | | | 0.043 | |

¹Based on linear quadratic model (dose+dose²).

Table 4

Comparison of *RET/PTC* or *PAX8/PPAR γ* positive tumors with *BRAF* or *RAS* mutation positive tumors according to selected patient characteristics in thyroid cancers that developed after the Chernobyl accident

| Characteristic | Relative risk of rearrangements vs point mutations (95% CI) | | P heterogeneity ¹ |
|---------------------|--|------------------------|------------------------------|
| I-131 dose, Gy | | 8.00 (1.30, 150.96) | 0.020 |
| Oblast of residence | Kyiv vs Chernihiv | 4.12 (0.22, 100.89) | 0.003 |
| | Zhytomyr vs Chernihiv | 139.07 (7.13, 5843.00) | |
| Age at surgery, yr | | 0.62 (0.47, 0.79) | <0.001 |
| Sex | Female vs Male | 0.01 (0.00, 0.16) | <0.001 |

¹Based on multivariate logistic regression models.