

Optimal Detection of Sentinel Lymph Node Metastases by Intraoperative Radioactive Threshold and Molecular Analysis in Patients with Melanoma

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The aim of this study was to optimize a protocol for radioguided biopsy of the sentinel lymph node (SLN) in patients with melanoma. The protocol was based on a combination of ex vivo counting of the nodes detected intraoperatively and analysis of the harvested nodes by hematoxylin and eosin staining plus immunohistochemistry (conventional histopathology [PATH]) and by molecular biology (reverse-transcriptase polymerase chain reaction [RT-PCR]). **Methods:** A total of 124 patients with primary clinical stage I–II (according to the American Joint Committee on Cancer) cutaneous melanoma underwent successful radioguided SLN biopsy. SLNs harvested for analysis included any additional nodes whose ex vivo counting rate exceeded 20% of the hottest node. All removed SLNs were examined by conventional PATH and with RT-PCR analysis for the expression of messenger RNA for tyrosinase and the melanoma antigens recognized by T cells. Complete lymph node dissection (CLND) was performed only in the case of SLN metastasis detected by PATH. Different combinations of the intraoperative parameters (only the hottest node and all nodes harvested) and of analysis (PATH and RT-PCR) were tested as predictors of clinical outcome on the basis of long-term follow-up (12–81 mo; median, 55 mo). **Results:** A total of 197 SLNs were harvested, 41 of which harbored metastasis as detected by RT-PCR analysis; PATH detected metastasis in only 24 of 41 metastatic SLNs. In 5 of 41 instances, metastasis was not in the hottest SLN. The main factor determining correct classification of the SLN status was RT-PCR, which significantly improved detection of metastasis, even if applied only to the hottest node ($P < 0.0001$ vs. PATH analysis of either the hottest SLN or all nodes above the 20% threshold). Metastatic disease recurred locally in 5 patients who had not undergone CLND; RT-PCR analysis showed metastasis in 4 of these patients. The false-negative rate of SLN biopsy progressively decreased when applying PATH only to the hottest node (32.1%), additional RT-PCR to the hottest node (21.4%), PATH to all nodes (17.9%), and RT-PCR to all nodes (3.6%, $P = 0.015$ vs. PATH analysis of

only the hottest SLN). **Conclusion:** On the basis of long-term follow-up (the gold standard for final clinical outcome of SLN biopsy), both 20% threshold and RT-PCR analysis should be applied for optimal detection of nodal metastases in patients with melanoma.

Key Words: malignant cutaneous melanoma; radioguided sentinel lymph node biopsy; intraoperative ex-vivo counting; radioactive threshold; molecular analysis; clinical outcome

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Lymphatic mapping and sentinel lymph node (SLN) biopsy have rapidly emerged as the standard of care for patients with malignant cutaneous melanoma, as the procedure readily identifies those patients who may benefit from immediate complete lymph node dissection (CLND) of the lymphatic basin at risk for metastasis (1,2). In fact, the disease-free survival rate is significantly greater when CLND is performed immediately after a positive SLN biopsy than when nodal metastasis becomes clinically evident (78.5% vs. 73%, $P = 0.009$, at a median follow-up of 59.8 mo) (3).

Definition of the SLN as a biologic rather than an anatomic entity (i.e., not simply the node that lies closer to the tumor, but the regional node to which a tumor would first metastasize (4)) suggests some technical difficulties in the identification of the true SLN in the surgical field. In fact, multiple radioactive lymph nodes in the same lymphatic basin are often detected by lymphoscintigraphy or by intraoperative γ -probe counting. Especially if the progress of the radiocolloid has not been adequately monitored during lymphoscintigraphy, it is not always clear whether all these radioactive lymph nodes represent true SLNs or whether they simply are second-echelon nodes sequentially visualized by radiocolloid particles passing through the first, true SLN.

Therefore, different approaches for performing radioguided SLN biopsy have been described on the basis of the level of

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radioactivity detected within the nodes, either intraoperatively or by immediate ex vivo counting. Some authors base SLN identification on the absolute number of counts in the nodes, whereas others consider the ratio of the in vivo or ex vivo radioactive counts in the node relative to background or to neighboring non-SLNs (5,6). All such modalities are somewhat arbitrary, and debate is still open regarding the optimal strategy for removing radioactive lymph nodes for pathologic examination. An empiric threshold corresponding to 10% or more of the counting rate in the hottest SLN is widely reported in the literature (7), but it may lead to the superfluous removal of multiple non-SLNs (4).

The false-negative rate of SLN biopsy (i.e., the proportion of patients in whom metastatic disease appears during follow-up in a lymphatic basin that had been classified as disease-free on the basis of a negative SLN biopsy) ranges from 10% to 15% in patients with melanoma (8). Part of these false-negative cases is due to inadequate histopathologic analysis, such as hematoxylin and eosin (H&E) staining alone (9). Immunohistochemistry (IHC), and especially molecular biology techniques (e.g., reverse-transcriptase polymerase chain reaction [RT-PCR]), improve dramatically the capability of detecting microscopic metastatic disease in SLNs of patients with melanoma. Nevertheless, although IHC with antibodies to the S-100 (10) or HMB-45 antigens (11) has 10%–30% higher sensitivity for identifying micro-metastases than does H&E staining (9), it still has a sizable false-negative rate (6%–11.5%) (12–14). On the other hand, the RT-PCR technique upstages an additional 13%–30% of patients in whom SLNs were negative when analyzed by both H&E and IHC staining (15–18).

In this work, we define a protocol for radioguided SLN biopsy in patients with melanoma—a protocol that minimizes the number of radioactive lymph nodes removed and at the same time minimizes the risk of missing microscopic metastatic disease in the nodes removed. In particular, our goal was to verify that combining removal of radioactive lymph nodes that were 20% or more of the ex vivo counting rate in the hottest lymph node with molecular analysis in addition to H&E and IHC staining represents the best compromise in this setting. Long-term follow-up represented the gold standard for final evaluation of the clinical outcome of the procedure.

MATERIALS AND METHODS

Patients

The study includes 124 consecutive patients with cutaneous melanoma who underwent successful radioguided SLN biopsy. Informed written consent was obtained for the protocol, which was approved by the Ethical Committee of the University Hospital of Pisa. The primary melanomas were either a Breslow thickness of less than 1 mm and a Clark level III–V (clinical stage I and II, according to the American Joint Committee on Cancer) (19) or a Breslow thickness of 1 mm or more (stage II), with or without ulceration.

[Table 1] Patient characteristics are summarized in Table 1. According to current established guidelines, clinical decisions for further surgical or other medical treatment were made solely on the basis of PATH

TABLE 1
Clinical Characteristics of the 124 Patients Included in Study

Characteristic	Value
Median age (y)	56 (range, 20–81)
Sex (%)	
Male	53.20
Female	46.80
Primary tumor site (%)	
Nonextremities	56.60
Extremities	43.40
Histology (%)	
Superficial spreading	53.60
Other types	46.40
Breslow thickness	
Mean \pm SD (mm)	2.0 \pm 1.18
Median (mm)	1.6 (range, 0.4–6.0)
<2.0 mm (%)	62.70
2.01–4.0 mm (%)	31.70
>4.0 mm (%)	5.60
Clark stage (%)	
I–III	34.60
IV–V	65.40
Ulceration (%)	
Present	14.20
Absent	85.80
American Joint Committee on Cancer clinical stage (%)	
IA	11.90
IB	46.80
IIA	31.00
IIB	10.30

analysis (20). Thus, 23 patients in whom SLNs were metastatic by H&E and IHC (conventional histopathology [PATH]) underwent CLND of the affected basin, and 19 of these 23 patients were offered adjuvant therapy with either high-dose (13 patients) or low-dose (6 patients) interferon- α -2b (IFN, Intron-A; Schering-Plough) (21). However, no lymphadenectomy or adjuvant IFN therapy was offered to patients in whom SLNs were negative by PATH, even in the presence of a positive result at molecular analysis. The follow-up schedule after primary treatment or staging was reported by Romanini et al. (18). Median follow-up for the entire patient population was 54.8 mo (range, 11.6–80.6 mo).

Lymphatic Mapping and Radioguided SLN Biopsy

Lymphoscintigraphy was performed 6–18 h before SLN biopsy as described earlier (18). Briefly, 0.1- to 0.2-mL aliquots containing 4–8 MBq of ^{99m}Tc -albumin nanocolloid (Nanocol; GE Healthcare) were injected intradermally around the primary melanoma or on skin margins of the surgical scar if the primary lesion had previously been excised (22). Both early dynamic imaging (1 frame/min for 30 min) and delayed static imaging (various projections, as appropriate) were recorded using a large-field-of-view γ -camera (Camstar XRT; GE Healthcare) equipped with a parallel-hole, high-resolution collimator ($\pm 10\%$ window centered on the 140-keV energy peak of ^{99m}Tc). The skin projection of the SLN was confirmed by external counting with the γ -probe and marked with indelible ink (22).

SLN biopsy was performed under intraoperative radioguidance (22) using a handheld, thallium-doped cesium iodide-based γ -probe (Scintiprobe MR 100; Pol.hi.tech). Both the hottest lymph node and any additional radioactive lymph node counting 20% or more of the

ex vivo counting rate of the hottest node were harvested for analysis. Removed lymph nodes were labeled as the hottest node or as an additional hot node, as appropriate. When the radiocolloid drained into more than 1 nodal basin, SLN biopsy was performed for each draining basin.

PATH and RT-PCR Analysis

The harvested SLNs were immediately bisected; one half of each SLN was sent for PATH analysis and the other half was placed in Trizol (Life Technology) for storage at -80°C and delayed RT-PCR molecular analysis.

Details of PATH analysis have been described previously (18). Briefly, 5- μm -thick sections were cut from formalin-fixed, paraffin-embedded blocks. If H&E staining revealed metastasis, no further staining was performed. If H&E staining was negative for metastasis, adjacent sections were processed for IHC using antibodies for the S-100 and HMB-45 proteins; further staining with antibody to the *MART-1* antigen was performed in case of negative S-100 and HMB-45 staining (23). The entire sections were examined with 40 \times and 100 \times magnifications to detect discrete foci of tumor cells. The sinusoids were examined at 250 \times magnification to detect small clumps or single tumor cells.

For molecular analysis, total RNA was extracted from homogenized tissues and stored in 2- μg aliquots under ethanol at -80°C until tested. One microgram of RNA was reverse-transcribed into cDNA in a 20- μL reaction mixture containing 100 pmol of random hexamers (Roche), 6 units of reverse transcriptase (Promega), 6.0 mM magnesium chloride, 0.2 mM deoxynucleotide triphosphate, and 8 units of RNase inhibitor (Invitrogen). Samples were sequentially incubated at 25°C (10 min), 42°C (60 min), and 95°C (5 min) to inactivate the reverse transcriptase. cDNA samples were subsequently amplified for the target sequence by using the following reaction mixture: 5 μL of cDNA, 1.5 mM magnesium chloride, 0.2 mM deoxynucleotide triphosphate, 1 unit of AmpliTaq Gold DNA polymerase (Applied Biosystems), 20 pmol of each primer, and PCR buffer 1 \times in 25 μL of final volume. Each sample was first tested for amplifiability with glyceraldehydes-3-phosphate dehydrogenase primers (yielding a 320-base pair [bp] amplicon (24)) and then analyzed for the melanoma-associated markers *Tyr* and *MART-1* (generating 207- and 364-bp amplicons, respectively (25,26)). An ABI 2400 thermal cycler (Thermo Fisher Scientific) was used for amplification, applying the cycling profile described previously (27). After the reaction was complete, 20 μL of PCR samples were run on 2% agarose gel and scored positively when amplicons of the expected size were detected under ultraviolet light.

To minimize the risk of cross-contaminations, RNA extraction, setup of reaction mixtures, and amplifications were performed in separated rooms, and the SLN samples were intermingled with no template samples (negative controls) and RNA extracted from the SK-mel 23 melanoma cell line (positive control).

Classification of SLN by Different Criteria

Four sets of criteria for classifying SLNs as metastatic or non-metastatic were derived by variously combining the intraoperative data (ex vivo counting relative to the 20% threshold) with the results of SLN analysis. The most sensitive indicator available—molecular analysis—was taken as the reference standard for the presence of metastasis in the lymph nodes removed. The following combinations were identified: PATH analysis of only the hottest node, regardless of RT-PCR analysis (scenario A); PATH analysis of all nodes removed, regardless of RT-PCR analysis (scenario B); PATH

and PCR analysis of only the hottest node (scenario C); and PATH and PCR analysis of all nodes removed (scenario D).

False-Negative Rate

The false-negative rate was calculated as the number of false-negative cases over the sum of the true-positive plus the false-negative cases (28). True-positive cases were those patients who underwent CLND and in whom the SLN was classified as metastatic by PATH and those patients who did not undergo CLND but were classified as positive by RT-PCR analysis and, during long-term follow-up, presented with locoregional recurrence in a lymphatic basin that had been classified as metastasis-free by PATH analysis. False-negative cases were those patients who, during long-term follow-up, presented with locoregional recurrence in a lymphatic basin that had been classified as metastasis-free by either PATH or RT-PCR analysis. Patients exhibiting distant or in-transit metastasis (without lymph node metastasis in the lymphatic basin of interest) were not considered false-negative cases. False-negative rate was calculated for each of 4 classification scenarios.

Statistical Analysis

Nonparametric tests were used to compare proportions of reliable SLN biopsies defined according to the 4 different classification scenarios. Data were analyzed using the SPSS/PC+ 11.5 statistical software (SPSS Inc.), considering a *P* value of less than 0.05 as significant.

RESULTS

Classification of SLNs

A total of 197 SLNs were identified by intraoperative γ -probe counting (84.8% of which were detected only by preoperative lymphoscintigraphy) in 152 draining nodal basins of the 124 patients (1.23 basins per patient, 1.3 SLNs per basin). A total of 131 SLNs in 85 patients did not have metastasis when examined by both PATH and RT-PCR analysis; 24 of 44 harvested SLNs in 23 patients were classified as metastatic by both PATH and RT-PCR analysis; and 17 of 22 harvested SLNs in 16 patients were classified as metastatic by RT-PCR analysis only. No instance in which an RT-PCR-negative SLN was metastatic by PATH analysis was demonstrated, and no significant difference was found in the ex vivo counting rate of negative versus metastatic SLNs (data not shown).

A single SLN was found in 121 of the total 152 draining lymphatic basins, whereas 31 basins had multiple SLNs (a total of 76 SLNs; mean, 2.45 SLNs per basin; range, 2–4 SLNs per basin). Fifteen of 31 lymphatic basins from which more than 1 SLN were harvested harbored metastasis. In 10 of 15 of these tumor-positive basins, metastasis was found in the hottest lymph node, whereas in 5 cases the metastatic lymph node was less radioactive than the hottest node (Fig. 1). [Fig. 1] Thus, in 5 of 31 basins with multiple SLNs, metastasis was found in a node other than the hottest one (16.1%). Table 2 [Table 2] shows the location, ex vivo counting rate, and PATH/PCR analysis of these 5 metastatic nodes. Although in 4 of 5 of these nodes metastasis was detected both by PATH and by molecular analysis, it was only RT-PCR that detected metastasis in the remaining node (located in the parotid gland).

Table 3 reports the rates of correct classification for the 41 [Table 3] metastatic SLNs, as derived from the 4 classification scenarios

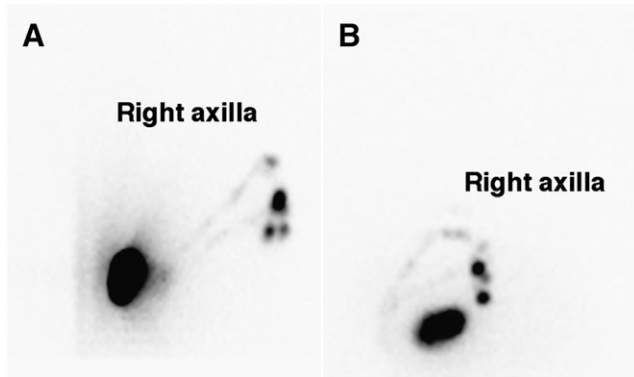


FIGURE 1. Lymphoscintigram (lateral [A] and oblique anterior [B] projections) obtained from patient who underwent surgery for cutaneous melanoma on her back. Lymph flowed through at least 2 main channels, converging at 4 radioactive nodes in right axilla. After intraoperative 20% threshold, 4 sentinel nodes were removed surgically. Ex vivo counting rate of third SLN removed (positive for tumor by both PATH and RT-PCR analyses) was equal to 35% of hottest sentinel node, which, like remaining 2, was negative. This is case in 1 of 5 patients in whom presence of metastasis would have been missed if only hottest node had been harvested for analysis.

(A, B, C, and D). The main determinant of correct classification was molecular analysis, which significantly increased the correct classification rate even if applied to only the hottest node, from 48.8% of combination A to 87.8% of combination C ($P < 0.0001$). On the other hand, when PATH analysis only was considered, removing additional radioactive lymph nodes did not significantly increase the correct classification rate for analyzing only the hottest node (from 48.8% of combination A to 58.5% of combination B, $P = 0.5$).

Clinical Outcome

Per current guidelines, CLND was performed only on those 23 patients in whom SLN metastasis was detected by PATH analysis; thus, 16 patients in whom metastasis was detected only by RT-PCR analysis did not undergo CLND of the affected basin. The clinical outcome of the entire patient population followed for a median of 55 mo (range, 12–81 mo) is charted in Figure 2. During long-term follow-up, none of the 23 patients who underwent CLND had locoregional tumor

recurrence (0%), whereas distant metastasis developed in 15 of these 23 patients and 11 of them died because of such systemic recurrences. By contrast, locoregional recurrence developed in 4 of 16 patients (or 25%) who had not undergone CLND because metastasis had been detected only by RT-PCR analysis, whereas 2 of these patients had systemic recurrences. Three of 16 patients died in this group (2 because of metastatic disease and 1 because of local recurrence that subsequently developed into systemic disease). For the 85 patients in whom SLNs were negative by both PATH and RT-PCR analyses, there was only 1 case of locoregional recurrence (1.2%), whereas distant metastasis developed in 12 of 85 patients during long-term follow-up, and 9 died.

False-Negative Rate of SLN Biopsy

A total of 5 locoregional recurrences were observed during follow-up, none of which was in the patients who underwent CLND. All 5 locoregional recurrences were false-negative on PATH analysis, whereas only 1 was false-negative on RT-PCR analysis. The contribution of each set of data to the false-negative rate of SLN biopsy was assessed by modeling different scenarios based on classification of the SLN status according to the 4 classification scenarios (Table 4). The worst scenario was that using the least-sensitive technique of analysis (PATH) to analyze only the hottest lymph node (scenario A). In this case, the false-negative rate of SLN biopsy was 9 of 28, or 32.1%. Including in the PATH analysis all lymph nodes harvested above the 20% ex vivo counting rate of the hottest node (scenario B) reduced the false-negative rate of SLN biopsy to 5 of 28, or 17.9%. The false-negative rate estimated when using the most-sensitive technique of SLN analysis (RT-PCR) (scenario C), compared with the hottest node only, was 6 of 28, or 21.4%, whereas the greatest reduction in the false-negative rate was expected when using RT-PCR analysis for all nodes harvested (scenario D) (1 of 28, or 3.6%). There was a statistically significant difference only for scenario D versus scenario A ($P = 0.015$).

DISCUSSION

The SLN status represents a highly significant prognostic factor in patients with clinically localized cutaneous mela-

TABLE 2
Metastases in Nodes Other Than That with Highest Radioactivity

Lymphatic basin	Hottest node*	Metastatic node*	Relative to hottest node (%)	Order of harvesting	Status
Axilla	1,356	1,200	88.40	Second of 3	PATH+/RT-PCR+
Parotid	68	34	50.00	Second of 2	PATH-/RT-PCR+
Axilla	832	322	38.70	Second of 3	PATH+/RT-PCR+
Axilla	455	160	35.10	Third of 4	PATH+/RT-PCR+
Axilla	900	300	33.30	Second of 3	PATH+/RT-PCR+

*Ex vivo counting rate measured intraoperatively with handheld γ -probe; counts/10 s.

PATH+ = PATH-positive; PCR+ = PCR-positive.

In these patients, presence of metastasis would have been missed if only hottest node had been harvested for analysis.

TABLE 3
Rates of Correct Classification of Metastatic SLNs

Classification	Number correctly classified	Correct classification rate (%)
A	20/41	48.80
B	24/41	58.50
C	36/41	87.80
D	41/41*	100.0*

*Standard of reference.

B vs. A, $P = 0.5$; C vs. B, $P = 0.005$; C vs. A, $P < 0.0001$; D vs. C, $P = 0.06$; D vs. B, $P < 0.0001$; D vs. A, $P \ll 0.0001$.

noma (3,19). On average, the SLN is metastatic by conventional PATH analysis in 1 of 5 or 6 patients with stage I–II melanoma. Among the same population with a negative SLN, locoregional recurrence still occurs during follow-up in up to 15% of cases, which presumably represent failures of nuclear medicine, surgery, or histopathology (8). Although radio-guided SLN biopsy has rapidly gained the favor of surgeons, the optimal criteria are still debated both for intraoperative SLN identification and for detection of micrometastasis. Although initial spread of melanoma to regional lymph nodes follows, in general, an orderly pattern, separate parallel draining to more than 1 lymph node (occasionally located in different lymphatic basins) is frequently observed. Routine lymphoscintigraphy is, therefore, important in identifying unsuspected locations of draining lymph nodes, even in patients with melanomas located in the extremities (29).

Radioactive SLNs can readily be identified intraoperatively because of their much higher counting rates relative to background. When histologically positive SLNs are found, they are usually, but not always, the hottest nodes. Several

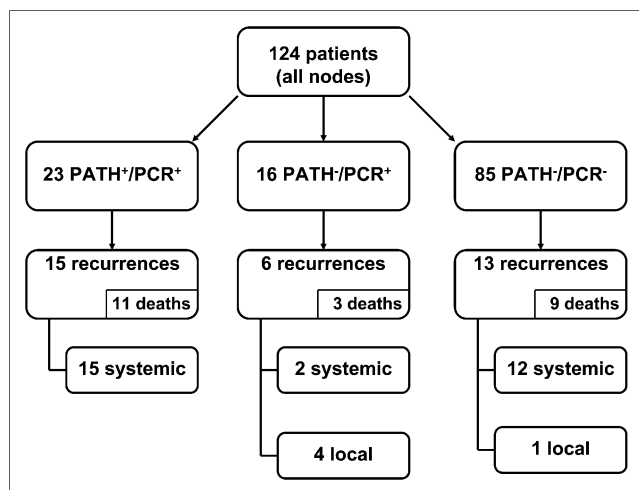


FIGURE 2. Clinical outcome observed in follow-up as function of SLN status evaluated by PATH or by RT-PCR analysis of molecular markers. PATH– = conventional histopathology negative; PATH+ = conventional histopathology positive; PCR– = RT-PCR negative; PCR+ = RT-PCR positive.

TABLE 4
Results for the 4 Classification Scenarios

Parameter	Scenario			
	A	B	C	D
FN cases (n)	5* + 4†	5*	2* + 4†	1*
TP cases (n)	19‡	23‡	19‡ + 3§	23‡ + 4§
FN rate	9/(9 + 19)	5/(5 + 23)	6/(6 + 22)	1/(1 + 27)
FN rate (%)	32.14	17.86	21.43	3.57

*Local recurrences observed.

†Additional false-negative cases, which would have been potentially observed if only hottest node had been harvested.

‡Patients with SLN classified as metastatic by PATH.

§Patients PATH negative and PCR positive with nodal recurrence during long-term follow-up.

FN = false-negative; TP = true-positive.

FN rate = FN/(FN + TP).

FN rate (%) = (FN/[FN + TP]) × 100.

operational definitions of the SLN have evolved over time to help the surgeon decide exactly which nodes should be removed to maximize the likelihood of locating the true biologic SLN and minimize the superfluous removal of multiple non-SLNs (30–35). An operational SLN definition that was associated with the lowest false-negative rate, using the extensive database of the Sunbelt Melanoma Trial, was described as “. . . all nodes containing radioactive counts ≥ 10% that of the hottest nodes removed” (7,36). However, these authors did not validate such an operational definition with, for example, long-term follow-up for nodal recurrence in basins found to be free from metastasis at SLN biopsy.

Because the 10% threshold could lead to the superfluous removal of multiple non-SLNs, Coit suggested that such a threshold could be increased to, for example, 20% or even 30% (4). Among those thresholds suggested by Coit, for this study we adopted the threshold closest to that proposed by McMasters et al. (i.e., 20% rather than 10%) (7) as a new operational definition of SLN. We validated the reliability of this procedure with long-term follow-up for identifying the false-negative cases.

Detection of micrometastatic disease is an additional crucial determinant of the false-negative rate for SLN biopsy in patients with melanoma. In this regard, it is now being increasingly accepted that, despite some controversies still surrounding this issue (37), molecular biology techniques such as RT-PCR support an improvement in sensitivity that seems to come at the cost of increased false-positives and, thus, diminished specificity when compared with histologic analysis. False-positives in tyrosinase mRNA analysis may arise from assay contamination, illegitimate transcription, or legitimate sources such as tyrosinase-expressing Schwann cells and benign nevus cells (15–18). A recent metaanalysis of 22 studies totaling 4,019 patients with clinical stage I or II melanoma who underwent SLN biopsy (38) concluded that detecting SLN metastasis by RT-PCR correlates significantly with the likelihood of disease recurrence. In particular, the

hazard ratio for disease-free survival and for overall survival is 5-fold and 3-fold, respectively, higher for the patients who were RT-PCR-positive than for those who were RT-PCR-negative. In our study, RT-PCR analysis identified an additional 17 metastatic SLNs over the 24 SLNs that PATH analysis had identified as metastatic.

By combining the ex vivo counting rate with PATH plus RT-PCR analysis, we tested 4 different combinations of SLN status and found that molecular analysis is the most crucial factor for correct classification of SLN status because RT-PCR statistically improves the correct classification rate, even when applied to only the hottest SLN.

The long-term follow-up we adopted (median, 55 mo) overcomes a potential bias of our study, that is, that we did not harvest for analysis radioactive lymph nodes with counting rates higher than the 10% threshold proposed by McMasters et al. (7) but lower than the 20% threshold adopted in this work. In our series, even a 30% counting rate threshold would have been sufficient for accurately defining the SLN status with PATH analysis, which is currently the recommended operational standard for selecting patients for CLND (Table 4).

Five patients had locoregional recurrences during long-term follow-up; all of these were false-negative cases on PATH, but only 1 was false-negative on RT-PCR analysis. Thus, adding RT-PCR to PATH analysis of all SLNs above the 20% threshold of the hottest node results in an absolute reduction of the false-negative rate from 17.9% to 3.6%. On the other hand, if we had removed only the hottest node and used only PATH for analysis (scenario A), we would have potentially observed 4 additional false-negative cases. Added to the 5 local recurrences actually observed, this would have resulted in a 32.1% false-negative rate, significantly greater ($P = 0.015$) than the value we would expect when analyzing all SLNs with RT-PCR (scenario D). Taken altogether, these observations underscore the importance of not only RT-PCR analysis for a correct classification of the SLN(s) but also the harvesting of all nodes with counting rates above the 20% threshold relative to the hottest node. Only 1 SLN was detected by lymphoscintigraphy and removed in the single patient who presented locoregional recurrence of the 85 PATH- and PCR-negative cases. This observation suggests a failure of lymphoscintigraphy or RT-PCR analysis or disease in transit.

The results of the Multicenter Selective Lymphadenectomy Trial-I report that SLN status has a significant prognostic power, and lymphadenectomy after detection of clinically occult SLN metastases significantly prolongs disease-free survival, compared with therapeutic lymphadenectomy when SLNs grow to a palpable size (3). Uncertainty still remains as to how to treat PATH-negative and PCR-positive cases. This issue is being addressed by the second Multicenter Selective Lymphadenectomy Trial, designed to evaluate the effect of regional lymphadenectomy on disease-free and overall survival in patients with early nodal metastases detected by RT-PCR analysis (39).

CONCLUSION

International guidelines lack a unique definition for the surgical detection of SLNs, possibly contributing to the relatively high false-negative rate of SLN biopsy (28). It is, therefore, mandatory for the nuclear medicine community to reach a consensus on the radioactive counting rate threshold to best guide the surgeon in the identification of those lymph nodes with the highest probability of harboring metastases (true biologic SLNs), thus avoiding the unnecessary removal of radioactive non-SLNs and reducing the morbidity related to the procedure. The results of the present study will help the nuclear medicine community to reach such a consensus.

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