

Sinchita Roy-Chowdhuri
Paul A. VanderLaan · John M. Stewart
Gilda da Cunha Santos *Editors*



Molecular Diagnostics in Cytopathology

A Practical Handbook for
the Practicing Pathologist



Springer

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Preface

This book is a compilation of high-yield molecular pathology topics relevant to the field of cytopathology in a user-friendly format that can be used as a reference handbook by training and practicing pathologists and laboratory personnel dealing with, and interested in, this evolving field.

Molecular diagnostics are increasingly used to help guide targeted therapy in solid organ tumors and hematologic malignancies. A large proportion of molecular testing is performed on limited-volume samples obtained via minimally invasive techniques, such as fine needle aspiration. Increasingly, cytopathologists play an essential role in this process, both in the triage of specimens during rapid on-site evaluation and in the evaluation of archival samples to determine suitability for ancillary testing. Therefore, it is imperative that practicing cytopathologists stay abreast of up-to-date diagnostic, prognostic, and predictive ancillary tests that can be used on limited cytologic material. This is a challenge since the landscape of known genomic alterations is constantly evolving and the subsequent set of testing options is ever expanding. In addition, many practicing cytopathologists have not had substantial molecular pathology training during residency or fellowship; therefore, the basic core principles of molecular testing may remain elusive.

The main focus of this book is to provide an overview of the principles of molecular diagnostics in context of cytopathology specimens together with a basic understanding and working knowledge of the available technology, platforms, and clinical applications. The initial sections of the book summarize the pre-analytic aspects of molecular testing, including

cytology specimen preparation and handling, specimen selection and evaluation, and workflow algorithms as well as the analytic considerations for a variety of nucleic acid and protein-based testing. The remaining section focuses on disease-specific molecular testing for various organ-based applications.

Molecular cytopathology is an expanding and evolving field. With the increasing demand for molecular testing on small specimens, cytopathologists need to understand how to efficiently select and triage the best tissue for testing as well as interpret the molecular test results in context of the morphology to cement the role of cytopathology as an independent and essential component of diagnostic and precision medicine.

We hope this book will serve as a practical handbook of clinical molecular diagnostics in context of the cytopathology specimen.

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Part I

Overview of Molecular Cytopathology



Chapter 1

Introduction: Overview of Current Molecular Diagnostic Testing on Cytology Samples

Michael H. Roh and Rashmi Kanagal-Shamanna

Abbreviations

<i>ALK</i>	Anaplastic lymphoma kinase or ALK receptor tyrosine kinase
<i>BRAF</i>	v-raf murine sarcoma viral oncogene homolog B
CEP	Centromeric probe
<i>DDIT3</i>	DNA damage-inducible transcript 3
<i>EGFR</i>	Epidermal growth factor receptor
<i>EWSR1</i>	Ewing sarcoma breakpoint region 1
FISH	Fluorescence <i>in situ</i> hybridization
FNA	Fine-needle aspiration
<i>FOXO1</i>	Forkhead box O1
<i>FUS</i>	Fused in sarcoma

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<i>GNAI1</i>	Guanine nucleotide-binding protein subunit alpha-11
<i>GNAQ</i>	Guanine nucleotide-binding protein G(q) subunit alpha
<i>HER2</i>	Human epidermal growth factor receptor 2 (<i>ERBB2</i>)
HPV	Human papillomavirus
<i>KIT</i>	KIT proto-oncogene receptor tyrosine kinase
<i>KRAS</i>	Kirsten rat sarcoma viral oncogene homolog
<i>MDM2</i>	Mouse double minute 2 homolog or E3 ubiquitin protein ligase homolog (mouse)
<i>MEK1</i>	Mitogen-activated protein kinase kinase 1
<i>MYCN</i>	v-myc myelocytomatosis viral oncogene homolog, neuroblastoma derived
<i>NRAS</i>	Neuroblastoma RAS viral oncogene
<i>PDGFB</i>	Platelet-derived growth factor β
<i>PDGFRA</i>	Platelet-derived growth factor receptor A
Pter	Terminal of chromosome short arm
Qter	Terminal of chromosome long arm
RNA	Ribonucleic acid
<i>ROS1</i>	ROS proto-oncogene 1, receptor tyrosine kinase
<i>SS18</i>	Synovial sarcoma translocation chromosome 18
<i>TFE3</i>	Transcription factor binding to immunoglobulin heavy constant

Key Points

- Molecular testing plays a critical role in the work-up of both solid tumors and hematological malignancies for the purposes of diagnosis and biomarker identification for assessment of prognosis and therapeutic targets
- Pathologists have a central role in integrating appropriate molecular ancillary testing into routine cytopathology workflow to enable personalized medicine

During this continuously evolving era of precision medicine, there have been tremendous advances in our understanding of the molecular genetic aberrations (e.g., mutations, gene amplifications, and gene rearrangements) that underlie and drive the growth and spread of various cancers. This has been complemented by advances in molecular diagnostic technologies available to pathologists and laboratory personnel to interrogate and detect these genetic abnormalities. Concurrent advances in minimally invasive interventional procedures have resulted in an increasing reliance on small biopsies along with exfoliative and fine-needle aspiration (FNA) cytology specimens not only for diagnostic purposes but also for ancillary molecular testing to guide the increasingly “personalized” management of patients with cancer [1–4]. In addition, the rapid development of new high-throughput and multiplex molecular testing modalities holds the promise to address the issue of simultaneously testing for a multitude of different genomic abnormalities utilizing cytology samples and small biopsies [5–8]. Furthermore, the recognition of tumor heterogeneity at the molecular level places an increasing demand for these procedures, as will the need to analyze multiple samples from the same patient over time in order to monitor the pattern of evolution of molecular abnormalities [9]. This would serve to reevaluate patients with cancers which have stopped responding to targeted therapy in order to identify new alterations which may be amenable to a switch in a targeted therapeutic regimen. Consequently, the importance of effectively integrating molecular ancillary testing with anatomic pathology workflow, particularly with regard to cytologic and small biopsy specimens, has never been greater.

Section II of this book will broadly address the practical considerations of cytology specimen processing in the molecular laboratory. Namely, chapters will be dedicated to the types of preparatory protocols and platforms utilized for molecular ancillary testing (Chap. 2), sample acquisition and test requisition (Chap. 3), specimen assessment/selection

along with triaging workflows for this purpose (Chap. 4), technical aspects of the various molecular diagnostic techniques and the practical implications for each (Chaps. 5, 6, and 7), and the interpretation of molecular diagnostic results along with how they correlate with cytopathologic diagnoses (Chap. 8).

Section III of this book will discuss the clinical settings in which molecular testing dovetails with the evaluation of cytology and small biopsy specimens, with respect to specific organ systems. In one aspect, molecular ancillary tests can be seen as adjuncts to aid in the diagnosis of small biopsies and cytology specimens [10–19]. Some key examples, not intended to be exhaustive, are listed in Table 1.1. The classic example is human papillomavirus (HPV) testing applied to gynecologic cytology specimens, which can be particularly useful to help guide the management of patients with indeterminate Pap test results [20]. HPV testing is also being increasingly applied in the diagnostic work-up of head and neck squamous cell carcinomas [21–24]. Detailed discussion of these applications will be addressed in Chaps. 9 and 10. Next, a variety of molecular diagnostic approaches have emerged to aid in the diagnosis of indeterminate thyroid FNAs [10, 15–17, 25–29]. These include high-throughput analysis of gene mutations and gene rearrangements, gene expression classifier analysis, and microRNA expression profiling (Chap. 12). Fluorescence *in situ* hybridization (FISH) analysis has also emerged as a useful diagnostic adjunct in the evaluation of pancreaticobiliary brushings (Chap. 15) and urinary tract cytology specimens (Chap. 16) [11, 12, 18]. FISH testing also serves an instrumental role in the diagnosis and subclassification of salivary gland neoplasms (Chap. 14) and sarcomas (Chap. 18) in FNA and small biopsy samples [14, 30, 31].

Alternatively, molecular diagnostic adjuncts can be utilized to interrogate biomarkers that provide information pertinent to prognosis and/or targeted therapeutic strategies [1, 32–43]. Salient examples are listed in Table 1.2 and include molecular testing of non-small cell lung carcinomas for *EGFR* and *BRAF* gene mutations along with *ALK* and *ROS1* gene rearrangements (Chap. 11); testing of metastatic

TABLE 1.1 Examples of molecular testing as diagnostic adjuncts applied to cytology and small biopsy specimens

Context	Molecular diagnostic adjuncts
Liquid-based cervical cytology	HPV testing
Head and neck squamous cell carcinoma	HPV testing
Thyroid FNA cytology	Gene mutation and rearrangement analysis Gene expression classifier testing microRNA analysis
Urine cytology	FISH probe set (CEP3, CEP7, CEP17, 9q21)
Pancreatobiliary brushing cytology	FISH probe set (CEP3, CEP7, CEP17, 9q21) FISH probe set (1q21, 7p12, 8q24, 9p21)
Renal cell carcinoma (examples below)	FISH probes
• Clear cell renal cell carcinoma	3pter/3qter
• Papillary renal cell carcinoma	CEP1, CEP7, CEP17
• Chromophobe renal cell carcinoma	CEP1, CEP7, CEP17
• Translocation-associated renal cell carcinoma	<i>TFE3</i>
Soft tissue tumors (examples below)	FISH probes and RT-PCR assays
• Alveolar rhabdomyosarcoma	<i>FOXO1</i>
• Alveolar soft part sarcoma	<i>TFE3</i>

(continued)

TABLE 1.1 (continued)

Context	Molecular diagnostic adjuncts
• Clear cell sarcoma	<i>EWSR1</i>
• Dermatofibrosarcoma protuberans	<i>PDGFB</i>
• Desmoplastic small round cell tumor	<i>EWSR1</i>
• Ewing sarcoma	<i>EWSR1</i>
• Myxoid/round cell liposarcoma	<i>DDIT3/FUS</i> , <i>DDIT3/EWSR1</i>
• Synovial sarcoma	<i>SS18</i>
• Well-differentiated liposarcoma	<i>MDM2</i>
Small B-cell lymphomas	
• Lymphoplasmacytic lymphoma	<i>MYD88</i> , <i>CXCR4</i> mutation analysis
• Follicular lymphoma	t(14;18)(q32;q21) <i>IGH/BCL2</i>
• Mantle cell lymphoma	t(11;14)(q13;q32) <i>CCND1/IGH</i>
• Hairy cell leukemia	<i>BRAF</i> mutation analysis
• Extranodal marginal zone lymphoma	t(11;18)(q21;q21) <i>BIRC3/MALT1</i> ; t(14;18)(q32;q21) <i>IGH/MALT1</i> ; t(3;14)(p14.1;q32) <i>FOXP1/IGH</i> ; t(1;14)(p22;q32) <i>BCL10/IGH</i>
Burkitt lymphoma	t(8;14)(q24;q32) <i>MYC/IGH</i>
Extranodal NK/T-cell lymphoma, nasal type	EBV testing
ALK-positive T-cell lymphoma	<i>ALK</i> rearrangement

TABLE 1.2 Examples of molecular testing as prognostic/theranostic adjuncts applied to cytology and small biopsy specimens

Context	Molecular diagnostic adjuncts
Non-small cell lung carcinoma	<i>EGFR</i> and <i>BRAF</i> mutation testing <i>ALK</i> and <i>ROS1</i> rearrangement testing PD-L1 testing
Melanoma	<i>BRAF</i> , <i>KIT</i> , <i>GNAQ</i> , <i>GNAI1</i> , <i>NRAS</i> , <i>MEK1</i> mutation testing
Colon adenocarcinoma	<i>KRAS</i> , <i>BRAF</i> mutation testing
Breast carcinoma	<i>ERBB2</i> amplification
Gastroesophageal adenocarcinoma	<i>ERBB2</i> amplification
Gastrointestinal stromal tumor	<i>KIT</i> and <i>PDGFRA</i> mutation testing
Neuroblastoma	<i>MYCN</i> amplification
Inflammatory myofibroblastic tumor	<i>ALK</i> rearrangement
Papillary thyroid carcinoma	<i>BRAF</i> mutation testing
Lymphoplasmacytic lymphoma	<i>MYD88</i> , <i>CXCR4</i> mutation testing
Diffuse large B-cell lymphoma	<i>MYC</i> rearrangement FISH testing Gene expression classifier testing Gene mutation analysis for <i>CARD11</i> , <i>CD79A</i> , <i>CD79B</i> , <i>EZH2</i> , <i>MYC</i> , <i>MYD88</i> , <i>TP53</i> , etc.
High-grade B-cell lymphoma	<i>MYC</i> , <i>BCL2</i> , <i>BCL6</i> rearrangement

(continued)

TABLE 1.2 (continued)

Context	Molecular diagnostic adjuncts
Chronic lymphocytic leukemia/small lymphocytic lymphoma	Deletions of <i>ATM</i> , <i>D13S319</i> , and <i>TP53</i> ; trisomy 12 Gene mutation analysis for <i>ATM</i> , <i>BIRC3</i> , <i>TP53</i> , <i>NOTCH1</i> , <i>SF3B1</i> , etc. Drug resistance mutation testing for <i>BTK</i> , <i>PLCG2</i> genes
Myeloid sarcomas	FISH testing for multiple rearrangements Gene mutation analysis including <i>ASXL1</i> , <i>CALR</i> , <i>DNMT3A</i> , <i>FLT3</i> , <i>IDH1/2</i> , <i>JAK2</i> , <i>KIT</i> , <i>MPL</i> , <i>NPM1</i> , <i>RUNX1</i> , <i>SF3B1</i> , <i>SRSF2</i> , <i>TET2</i> , <i>TP53</i> , etc.

melanomas for various mutations including *BRAF*, *NRAS*, and *KIT* gene mutations; *ERBB2* gene amplification testing for breast (Chap. 13) and gastroesophageal carcinomas; and a multitude of molecular testing for pediatric tumors (Chap. 19) for which *MYCN* gene amplification testing in neuroblastomas represents a classic example.

Similar to solid tumors, molecular testing is a critical adjunct for diagnosis and personalized therapy in hematological malignancies [44]. This is reflected in the recently revised 2017 WHO classification system for lymphoid and myeloid disorders where genetic results form a basis for diagnosis and classification in multiple disease groups [45, 46]. Cytology FNA is routinely used for work-up of tissue-based hematological malignancies including lymphomas and leukemias and provides ample opportunity for molecular testing [5, 47]. Typical scenarios are summarized below, while Chap. 17 will address these in greater detail.

Differentiation of low-grade lymphomas and reactive lymphoid proliferations is, by far, the most challenging task for pathologists. Molecular tools can be extremely helpful to identify clonal markers that can facilitate this distinction. These include detection of monoclonal VDJ rearrangements in immunoglobulin heavy chain (*IgH*), T-cell receptor (*TCR*)

beta or gamma receptor genes, as well as specific gene rearrangements and gene mutations in B-cell and T-cell lymphomas [47–56]. In the right clinical and morphologic context, FISH or PCR studies for characteristic gene rearrangements are helpful in sub-typing of small B-cell lymphomas [such as t(11;14)(q13;q32) *IGH/CCND1* in mantle cell lymphoma, t(14;18)(q32;q21) *IGH@-BCL2* in follicular lymphoma, etc.], certain large B-cell lymphomas such as Burkitt lymphoma [t(8;14)(q24;q32) *MYC-IGH@*], and *ALK*-positive T-cell lymphomas [t(2;5)(p23;q35)] [47, 51, 57, 58]. Assessment of translocations by PCR enables evaluation of measurable (minimal) residual disease [59]. Molecular testing is essential to identify biomarkers to assess prognosis and determine the choice of therapy. Some examples, which are in no way comprehensive, include molecular testing of lymphoplasmacytic lymphoma for mutations in *MYD88* and *CXCR4* [60]; testing of large B-cell lymphomas for *MYC* gene rearrangement as well as gene expression profile for distinguishing between germinal center and activated B-cell phenotype [61, 62]; testing of high-grade B-cell lymphomas for rearrangements and copy number changes in *MYC*, *BCL2*, and *BCL6* genes [63, 64]; testing of T-cell lymphomas for *ALK* gene rearrangement (58); and testing of myeloid sarcomas for recurrent genetic translocations such as t(15;17) and t(9;22) *BCR/ABL* and various gene mutations such as *FLT3* and *IDH1/2* [46, 65, 66]. Further, in the context of newly developed targeted therapies, testing for gene mutations that confer resistance to drugs such as *BTK* mutations in chronic lymphocytic leukemia/small lymphocytic lymphomas is emerging as a standard modality [67]. In most cases, mutation testing for multiple genes is best done using a multi-gene sequencing assay such as next-generation sequencing. Molecular testing for malignancy associated viruses, specifically Epstein-Barr virus, plays an important role in diagnosis and monitoring of a variety of hematological conditions including Hodgkin and non-Hodgkin lymphoma, immunodeficiency-associated lymphoproliferative disorders including post-stem cell transplant [68, 69]. These applications will be further elaborated in Chap. 17.

Although some molecular tests serve more as diagnostic aids (Table 1.1) and others serve more to provide prognostic/theranostic information (Table 1.2), as mentioned above, these two roles of molecular diagnostic tests applied to cytology specimens and small biopsies should not be seen as necessarily mutually exclusive. As the repertoire of targeted therapeutic regimens expands, based on our evolving scientific knowledge surrounding molecular biomarkers and targets, these two roles will become increasingly intertwined.

The objective of this book is several-fold. First, the advantages and limitations of cytology specimens will be discussed to assist in the understanding, on the part of pathologists and laboratory personnel, of how best to leverage these patient samples for diagnosis and molecular testing. Second, practical knowledge surrounding the acquisition and laboratory processing of these specimens, along with triage and workflow algorithms, will be shared. This information will be cemented by providing details surrounding the actual testing methodologies utilized for molecular diagnostic evaluation of these samples (Section II). Finally, clinically relevant contexts in which molecular diagnostics and cytopathology are becoming increasingly intertwined will be discussed (Section III).

It is the hope of the editors of this book and the individual chapter authors that the reader can utilize this book as a practical guide to develop best practice algorithms by which molecular testing is applied to the diagnosis and work-up of cytology and small biopsy specimens. Understanding the pre-analytic and analytic aspects of molecular testing for cytology samples will enable pathologists and laboratory personnel to best determine how samples are managed and triaged to successfully obtain the optimum amount of diagnostic, prognostic, and theranostic information for each patient sample.

References

1. Aisner DL, Sams SB. The role of cytology specimens in molecular testing of solid tumors: techniques, limitations, and opportunities. *Diagn Cytopathol.* 2012;40(6):511–24.
2. Clark DP. Seize the opportunity: underutilization of fine-needle aspiration biopsy to inform targeted cancer therapy decisions. *Cancer.* 2009;117(5):289–97.
3. Knoepp SM, Roh MH. Ancillary techniques on direct-smear aspirate slides: a significant evolution for cytopathology techniques. *Cancer Cytopathol.* 2013;121(3):120–8.
4. Krishnamurthy S. Applications of molecular techniques to fine-needle aspiration biopsy. *Cancer.* 2007;111(2):106–22.
5. Kanagal-Shamanna R, Portier BP, Singh RR, Routbort MJ, Aldape KD, Handal BA, et al. Next-generation sequencing-based multi-gene mutation profiling of solid tumors using fine needle aspiration samples: promises and challenges for routine clinical diagnostics. *Mod Pathol.* 2014;27(2):314–27.
6. Killian JK, Walker RL, Suuriniemi M, Jones L, Scurci S, Singh P, et al. Archival fine-needle aspiration cytopathology (FNAC) samples: untapped resource for clinical molecular profiling. *J Mol Diagn.* 2010;12(6):739–45.
7. Rekhtman N, Roy-Chowdhuri S. Cytology specimens: a goldmine for molecular testing. *Arch Pathol Lab Med.* 2016;140(11):1189–90.
8. Roy-Chowdhuri S, Chow CW, Kane MK, Yao H, Wistuba II, Krishnamurthy S, et al. Optimizing the DNA yield for molecular analysis from cytologic preparations. *Cancer Cytopathol.* 2016;124(4):254–60.
9. Beca F, Schmitt F. Growing indication for FNA to study and analyze tumor heterogeneity at metastatic sites. *Cancer Cytopathol.* 2014;122(7):504–11.
10. Alexander EK, Kennedy GC, Baloch ZW, Cibas ES, Chudova D, Diggans J, et al. Preoperative diagnosis of benign thyroid nodules with indeterminate cytology. *N Engl J Med.* 2012;367(8):705–15.
11. Barr Fritcher EG, Kipp BR, Halling KC, Clayton AC. FISHing for pancreatobiliary tract malignancy in endoscopic brushings

- enhances the sensitivity of routine cytology. *Cytopathology*. 2014;25(5):288–301.
12. Barr Fritcher EG, Voss JS, Brankley SM, Campion MB, Jenkins SM, Keeney ME, et al. An optimized set of fluorescence in situ hybridization probes for detection of pancreatobiliary tract cancer in cytology brush samples. *Gastroenterology*. 2015;149(7):1813–24. e1.
 13. Barroca H, Bom-Sucesso M. Fine needle biopsy with cytology in paediatrics: the importance of a multidisciplinary approach and the role of ancillary techniques. *Cytopathology*. 2014;25(1):6–20.
 14. Dal Cin P, Qian X, Cibas ES. The marriage of cytology and cytogenetics. *Cancer Cytopathol*. 2013;121(6):279–90.
 15. Nikiforov YE, Carty SE, Chiosea SI, Coyne C, Duvvuri U, Ferris RL, et al. Highly accurate diagnosis of cancer in thyroid nodules with follicular neoplasm/suspicious for a follicular neoplasm cytology by ThyroSeq v2 next-generation sequencing assay. *Cancer*. 2014;120(23):3627–34.
 16. Nikiforov YE, Carty SE, Chiosea SI, Coyne C, Duvvuri U, Ferris RL, et al. Impact of the multi-gene ThyroSeq next-generation sequencing assay on cancer diagnosis in thyroid nodules with atypia of undetermined significance/follicular lesion of undetermined significance cytology. *Thyroid*. 2015;25(11):1217–23.
 17. Nikiforov YE, Ohori NP, Hodak SP, Carty SE, LeBeau SO, Ferris RL, et al. Impact of mutational testing on the diagnosis and management of patients with cytologically indeterminate thyroid nodules: a prospective analysis of 1056 FNA samples. *J Clin Endocrinol Metab*. 2011;96(11):3390–7.
 18. Reynolds JP, Voss JS, Kipp BR, Karnes RJ, Nassar A, Clayton AC, et al. Comparison of urine cytology and fluorescence in situ hybridization in upper urothelial tract samples. *Cancer Cytopathol*. 2014;122(6):459–67.
 19. Savic S, Franco N, Grilli B, Barascud Ade V, Herzog M, Bode B, et al. Fluorescence in situ hybridization in the definitive diagnosis of malignant mesothelioma in effusion cytology. *Chest*. 2010;138(1):137–44.
 20. Massad LS, Einstein MH, Huh WK, Katki HA, Kinney WK, Schiffman M, et al. 2012 updated consensus guidelines for the management of abnormal cervical cancer screening tests and cancer precursors. *J Low Genit Tract Dis*. 2013;17(5 Suppl 1):S1–S27.
 21. Guo M, Khanna A, Dhillon J, Patel SJ, Feng J, Williams MD, et al. Cervista HPV assays for fine-needle aspiration

- specimens are a valid option for human papillomavirus testing in patients with oropharyngeal carcinoma. *Cancer Cytopathol.* 2014;122(2):96–103.
22. Guo M, Khanna A, Feng J, Patel S, Zhang W, Gong Y, et al. Analytical performance of cervista HPV 16/18 in SurePath pap specimens. *Diagn Cytopathol.* 2015;43(4):301–6.
 23. Kerr DA, Pitman MB, Sweeney B, Arpin RN 3rd, Wilbur DC, Faquin WC. Performance of the Roche cobas 4800 high-risk human papillomavirus test in cytologic preparations of squamous cell carcinoma of the head and neck. *Cancer Cytopathol.* 2014;122(3):167–74.
 24. Kerr DA, Sweeney B, Arpin RN 3rd, Ring M, Pitman MB, Wilbur DC, et al. Automated extraction of formalin-fixed, paraffin-embedded tissue for high-risk human papillomavirus testing of head and neck squamous cell carcinomas using the roche cobas 4800 system. *Arch Pathol Lab Med.* 2016;140(8):844–8.
 25. Agretti P, Ferrarini E, Rago T, Candelieri A, De Marco G, Dimida A, et al. MicroRNA expression profile helps to distinguish benign nodules from papillary thyroid carcinomas starting from cells of fine-needle aspiration. *Eur J Endocrinol.* 2012;167(3):393–400.
 26. Dettmer M, Vogetseder A, Durso MB, Moch H, Komminoth P, Perren A, et al. MicroRNA expression array identifies novel diagnostic markers for conventional and oncocytic follicular thyroid carcinomas. *J Clin Endocrinol Metab.* 2013;98(1):E1–7.
 27. Keutgen XM, Filicori F, Crowley MJ, Wang Y, Scognamiglio T, Hoda R, et al. A panel of four miRNAs accurately differentiates malignant from benign indeterminate thyroid lesions on fine needle aspiration. *Clin Cancer Res.* 2012;18(7):2032–8.
 28. Nikiforova MN, Tseng GC, Steward D, Diorio D, Nikiforov YE. MicroRNA expression profiling of thyroid tumors: biological significance and diagnostic utility. *J Clin Endocrinol Metab.* 2008;93(5):1600–8.
 29. Shen R, Liyanarachchi S, Li W, Wakely PE Jr, Saji M, Huang J, et al. MicroRNA signature in thyroid fine needle aspiration cytology applied to “atypia of undetermined significance” cases. *Thyroid.* 2012;22(1):9–16.
 30. Bajaj J, Gimenez C, Slim F, Aziz M, Das K. Fine-needle aspiration cytology of mammary analog secretory carcinoma masquerading as low-grade mucoepidermoid carcinoma: case report with a review of the literature. *Acta Cytol.* 2014;58(5):501–10.

31. Hudson JB, Collins BT. MYB gene abnormalities t(6;9) in adenoid cystic carcinoma fine-needle aspiration biopsy using fluorescence in situ hybridization. *Arch Pathol Lab Med.* 2014;138(3):403–9.
32. Aisner DL, Marshall CB. Molecular pathology of non-small cell lung cancer: a practical guide. *Am J Clin Pathol.* 2012;138(3):332–46.
33. Beltran H. The N-myc oncogene: maximizing its targets, regulation, and therapeutic potential. *Mol Cancer Res.* 2014;12(6):815–22.
34. Bergethson K, Shaw AT, Ou SH, Katayama R, Lovly CM, McDonald NT, et al. ROS1 rearrangements define a unique molecular class of lung cancers. *J Clin Oncol.* 2012;30(8):863–70.
35. Bernacki KD, Betz BL, Weigelin HC, Lao CD, Redman BG, Knoepp SM, et al. Molecular diagnostics of melanoma fine-needle aspirates: a cytology-histology correlation study. *Am J Clin Pathol.* 2012;138(5):670–7.
36. da Cunha Santos G, Saieg MA, Geddie W, Leighl N. EGFR gene status in cytological samples of nonsmall cell lung carcinoma: controversies and opportunities. *Cancer Cytopathol.* 2011;119(2):80–91.
37. Griewank KG, Scolyer RA, Thompson JF, Flaherty KT, Schadendorf D, Murali R. Genetic alterations and personalized medicine in melanoma: progress and future prospects. *J Natl Cancer Inst.* 2014;106(2):djt435.
38. Howell GM, Nikiforova MN, Carty SE, Armstrong MJ, Hodak SP, Stang MT, et al. BRAF V600E mutation independently predicts central compartment lymph node metastasis in patients with papillary thyroid cancer. *Ann Surg Oncol.* 2013;20(1):47–52.
39. Joensuu H, Hohenberger P, Corless CL. Gastrointestinal stromal tumour. *Lancet.* 2013;382(9896):973–83.
40. Pang NK, Nga ME, Chin SY, Ismail TM, Lim GL, Soong R, et al. KRAS and BRAF mutation analysis can be reliably performed on aspirated cytological specimens of metastatic colorectal carcinoma. *Cytopathology.* 2011;22(6):358–64.
41. Sipos JA, Shah MH. Thyroid cancer: emerging role for targeted therapies. *Ther Adv Med Oncol.* 2010;2(1):3–16.
42. Tothova Z, Wagner AJ. Anaplastic lymphoma kinase-directed therapy in inflammatory myofibroblastic tumors. *Curr Opin Oncol.* 2012;24(4):409–13.
43. Yoon HH, Sukov WR, Shi Q, Sattler CA, Wiktor AE, Diasio RB, et al. HER-2/neu gene amplification in relation to expression of

- HER2 and HER3 proteins in patients with esophageal adenocarcinoma. *Cancer*. 2014;120(3):415–24.
44. Kanagal-Shamanna R, Singh RR, Routbort MJ, Patel KP, Medeiros LJ, Luthra R. Principles of analytical validation of next-generation sequencing based mutational analysis for hematologic neoplasms in a CLIA-certified laboratory. *Expert Rev. Mol Diagn*. 2016;16(4):461–72.
 45. Swerdlow SH, Campo E, Pileri SA, Harris NL, Stein H, Siebert R, et al. The 2016 revision of the World Health Organization classification of lymphoid neoplasms. *Blood*. 2016;127(20):2375–90.
 46. Arber DA, Orazi A, Hasserjian R, Thiele J, Borowitz MJ, Le Beau MM, et al. The 2016 revision to the World Health Organization (WHO) classification of myeloid neoplasms and acute leukemia. *Blood*. 2016;127:2391–405; blood-2016-03-643544.
 47. Ochs RC, Bagg A. Molecular genetic characterization of lymphoma: application to cytology diagnosis. *Diagn Cytopathol*. 2012;40(6):542–55.
 48. Bagg A. Immunoglobulin and T-cell receptor gene rearrangements: minding your B's and T's in assessing lineage and clonality in neoplastic lymphoproliferative disorders. *J Mol Diagn*. 2006;8(4):426–9; quiz 526–7.
 49. Zhang S, Abreo F, Lowery-Nordberg M, Veillon DM, Cotelingam JD. The role of fluorescence in situ hybridization and polymerase chain reaction in the diagnosis and classification of lymphoproliferative disorders on fine-needle aspiration. *Cancer Cytopathol*. 2010;118(2):105–12.
 50. Maroto A, Martinez M, Martinez MA, de Agustin P, Rodriguez-Peralto JL. Comparative analysis of immunoglobulin polymerase chain reaction and flow cytometry in fine needle aspiration biopsy differential diagnosis of non-Hodgkin B-cell lymphoid malignancies. *Diagn Cytopathol*. 2009;37(9):647–53.
 51. Venkatraman L, Catherwood MA, Patterson A, Lioe TF, McCluggage WG, Anderson NH. Role of polymerase chain reaction and immunocytochemistry in the cytological assessment of lymphoid proliferations. *J Clin Pathol*. 2006;59(11):1160–5.
 52. Peluso AL, Ieni A, Mignogna C, Zeppa P. Lymph node fine-needle cytology: beyond flow cytometry. *Acta Cytol*. 2016;60(4):372–84.
 53. Grosso LE, Collins BT. DNA polymerase chain reaction using fine needle aspiration biopsy smears to evaluate non-Hodgkin's lymphoma. *Acta Cytol*. 1999;43(5):837–41.
 54. Langerak AW, Groenen PJ, Brüggemann M, Beldjord K, Bellan C, Bonello L, et al. EuroClonality/BIOMED-2 guidelines for

- interpretation and reporting of Ig/TCR clonality testing in suspected lymphoproliferations. *Leukemia*. 2012;26(10):2159.
55. Evans P, Pott C, Groenen P, Salles G, Davi F, Berger F, et al. Significantly improved PCR-based clonality testing in B-cell malignancies by use of multiple immunoglobulin gene targets. Report of the BIOMED-2 Concerted Action BHM4-CT98-3936. *Leukemia*. 2007;21(2):207.
 56. Patel KP, Pan Q, Wang Y, Maitta RW, Du J, Xue X, et al. Comparison of BIOMED-2 versus laboratory-developed polymerase chain reaction assays for detecting T-cell receptor- γ gene rearrangements. *J Mol Diagn*. 2010;12(2):226–37.
 57. Baro C, Espinet B, Salido M, Garcia M, Sanchez B, Florensa L, et al. Cryptic IGH/BCL2 rearrangements with variant FISH patterns in follicular lymphoma. *Leuk Res*. 2011;35(2):256–9.
 58. Shin HJ, Thorson P, Gu J, Katz RL. Detection of a subset of CD30+ anaplastic large cell lymphoma by interphase fluorescence in situ hybridization. *Diagn Cytopathol*. 2003;29(2):61–6.
 59. Rambaldi A, Carlotti E, Oldani E, Della Starza I, Baccarani M, Cortelazzo S, et al. Quantitative PCR of bone marrow BCL2/IgH+ cells at diagnosis predicts treatment response and long-term outcome in follicular non-Hodgkin lymphoma. *Blood*. 2005;105(9):3428–33.
 60. Treon SP, Cao Y, Xu L, Yang G, Liu X, Hunter ZR. Somatic mutations in MYD88 and CXCR4 are determinants of clinical presentation and overall survival in Waldenström macroglobulinemia. *Blood*. 2014;123(18):2791–6.
 61. Scott DW, Mottok A, Ennishi D, Wright GW, Farinha P, Ben-Neriah S, et al. Prognostic Significance of Diffuse Large B-Cell Lymphoma Cell of Origin Determined by Digital Gene Expression in Formalin-Fixed Paraffin-Embedded Tissue Biopsies. *J Clin Oncol*. 2015;33(26):2848–56.
 62. Landsburg DJ, Falkiewicz MK, Petrich AM, Chu BA, Behdad A, Li S, et al. Sole rearrangement but not amplification of MYC is associated with a poor prognosis in patients with diffuse large B cell lymphoma and B cell lymphoma unclassifiable. *Br J Haematol*. 2016;175(4):631–40.
 63. Li S, Desai P, Lin P, Yin CC, Tang G, Wang XJ, et al. MYC/BCL6 double-hit lymphoma (DHL): a tumour associated with an aggressive clinical course and poor prognosis. *Histopathology*. 2016;68(7):1090–8.