Molecular Diagnostics in Personalized Cancer Care

Best Practices and Overcoming Challenges



The content provided herein is for background and educational purposes only. The material is for your sole use and may not be altered or further disseminated in any fashion for further use.

PRECISION ONCOLOGY AND THE USE OF MOLECULAR BIOMARKERS EVOLVED FROM SCIENTIFIC BREAKTHROUGHS



CDx, companion diagnostic; ICI, immune checkpoint inhibitor; TCGA, The Cancer Genome Atlas.

Precision Medicine

Overview



Knowledge of tumor biology Targeted therapies

2017

First tumor-agnostic ICI using a genomic biomarker^{7,21}

2018

First tumor-agnostic targeted therapy^{2,3,22-24}

2018

TCGA published comprehensive list of driver mutations across 33 tumor types^{13,14}

Technological advances have contributed to **lower cost** and shorter turnaround time (TAT) for genome sequencing¹⁵⁻¹⁷

INCREASE OF THERAPEUTIC OPTIONS IN ONCOLOGY

Total Number of Anticancer Therapies Approved by the FDA Between 2009 and 2020²⁵



Between 2009 and 2020, there were 332 new anticancer therapy approvals, some of which require biomarker testing $^{\rm 25}$

As of June 2022, there are^{3,26}:

≥70 43 28 FDA-approved biomarkerlinked indications actionable genomic alterations by Precision Oncology

Biomarker testing is a fundamental component of precision oncology³

FDA, US Food and Drug Administration.



MOLECULAR DIAGNOSTICS IS A MULTISTEP PROCESS REQUIRING COLLABORATION AMONG DISTINCT DISCIPLINES^{27,28}



Communication and coordination between members of the core and expanded multidisciplinary team (MDT) are important to the implementation of precision oncology^{27,29,30}



DIAGNOSTIC JOURNEY IN PATIENTS WITH METASTATIC CANCER

Testing Navigation

The **Oncology nurse navigator** is a key point of contact between the patient and the MDT and aims to facilitate team communication and coordination during testing²⁷





SUCCESSFUL BIOMARKER TESTING DEPENDS ON KEY FACTORS



Testing **tissue** of sufficient quantity and quality³²



Use of appropriate tests³



Ordering process for actionable biomarkers³



Access to clear and searchable report data³³



TESTING FOR BIOMARKERS GENERALLY REQUIRES 20% OF TUMOR NUCLEI IN SAMPLES^{34,35}

Lung Adenocarcinoma Example

- Accurate detection of biomarkers may be difficult in samples with low numbers of tumor cells³⁶
- Interobserver variability and misestimation of tumor content are potential challenges^{34,36}
 - A study demonstrated that 38% of samples have overestimated tumor content³⁶
- Training may help lower discrepancies in estimating tumor content³⁴

Figure used with permission from Mikubo M et al. J Thorac Oncol. 2020;15(1):130-137.



Lung Adenocarcinoma Example³⁴ Tumor content **30%–40%**

Biopsy Choice May Impact Testing Outcomes



Biopsy Site

- Biomarker discordance between the primary tumor and a metastatic site may occur^{37,38}
- Additional/different drivers/ mutations may occur through clonal evolution over the course of the disease³⁹⁻⁴¹



Bone Biopsy

 Bone biopsy requires decalcification, which may impair sample yield and integrity, potentially negatively impacting biomarker testing outcomes⁴²



Rebiopsy

- Rebiopsy after disease progression may provide important and/or new information⁴³
- In certain cancers, receptor status may change over the course of the disease⁴⁴⁻⁴⁶



LIQUID BIOPSY OVERVIEW

Key Characteristics of Liquid Biopsy^{47,48}

ADVANTAGES

Is minimally invasive

Can capture tumor genetic heterogeneity and follow subclonal evolution through serial biopsy

Potentially represents genetic make-up from entire tumor and metastatic sites

May have a shorter overall TAT than tissue-based NGS relative to the date the test is ordered

Tissue Biopsy Testing

May provide a snapshot of the cellular and molecular characteristics of one part of a single tumor⁴⁹

Fresh tissue

 Does not provide information from all cancer cells

May miss an alteration if it is not present in the tested sample⁵⁰

Processing of biopsies of bone metastases may lead to DNA degradation⁵⁰

DISADVANTAGES

- Cannot directly correlate ctDNA results with histology or cellular phenotype
- Genetic analyses may have biased representation from differential tumor cell turnover

May be associated with false negatives

 Special processing and handling required

cfDNA/

ctDNA

Liquid Biopsy Testing

May reflect overall genomic landscape of the tumor and all metastatic sites (bone or other tissues)^{51,52}

• Does not provide information on TME⁵³

May miss an alteration if ctDNA concentration is below the LOD, leading to a false negative • ctDNA levels may vary

significantly^{52,54,55}

CTCs and ctDNA levels may be impacted by the number and sites of metastases, including bone^{52,54-56}

cfDNA, cell-free DNA; CTCs, circulating tumor cells; ctDNA, circulating tumor DNA; EVs, extracellular vesicles; FFPE, formalin-fixed, paraffin-embedded; LOD, limit of detection; TEPs, tumor-educated platelets; TME, tumor microenvironment. Image adapted from Alba-Bernal A et al. *EbioMedicine*. 2020;62:103100.



CHARACTERISTICS OF A GOOD BIOMARKER TEST

Clinical Guidelines and Expert Opinions⁵⁷⁻⁶³



Is actionable, prognostic, and/or predictive^{57,58}



Is supported by the highest level of evidence57



Has **tightly controlled specimen** collection, handling, and processing⁵⁷

Provides reproducible

results (>95%)^{59,60}



Has **predetermined cutoff** points/categories⁵⁷



Delivers **timely** results that impact treatment decisions⁶¹⁻⁶³



Possesses sufficient sensitivity, specificity, accuracy, and precision (<1% to 5% LOD) to detect actionable biomarkers⁵⁷⁻⁶⁰

ESSENTIAL QUESTIONS ABOUT A BIOMARKER TEST

(Analytical Validity	Clinical Validity	Clinical Utility
Definition ⁶⁴	The test is able to accurately and reliably measure the presence or absence of a biomarker in the appropriate specimen	The test can accurately and reliably identify a biologically defined disorder or separate into two or more groups with distinct clinical or biological outcomes or differences	The test has high levels of evidence that use of the biomarker can result in guiding clinical decisions that result in improved clinical outcomes compared with those if the biomarker test results were not applied
Essential question ⁴⁸	<i>Is the test for the biomarker sensitive, accurate, and reliable?</i>	Does the test accurately identify a disorder with distinct clinical or biological outcomes?	Is the test predictive of clinical outcomes?

Analytical and clinical validity is the foundation of *all* **biomarker testing.** In addition, to gain FDA approval, a CDx must be evaluated in a clinical study^{48,64-66}

Intended to depict biomarker testing methodologies. When testing for therapy selection, please consult product prescribing information and FDA-approved companion diagnostics.



BIOMARKER TESTING METHODS³



Images adapted with permission from Yu J et al. *Sci Rep.* 2019;9(1):7518, Yatabe Y et al. *J Thorac Oncol.* 2019;14(3):377-407, Kipf E et al. *J Mol Diagn.* 2022;24(1):57-68, and Goldbio. https://www.goldbio.com/articles/article/how-to-fragment-DNA-for-NGS. Accessed April 28, 2022.

USE OF APPROPRIATE TESTS

- Some biomarkers may be detected more reliably by **some specific testing technologies** than by others^{3,60}
- Gene rearrangements can be reliably detected by FISH and RNA-based NGS; enrichment strategy for a DNA-based NGS assay impacts the detection of fusions^{3,60}
- Understanding **assay limitations** is critical to identifying patients with actionable biomarkers³
- The American Society of Clinical Oncology (ASCO) recommends being familiar with the genomic testing platforms available to ensure fusion testing is performed when indicated³

Intended to depict biomarker testing methodologies. When testing for therapy selection, please consult product prescribing information and FDA-approved companion diagnostics.

NGS MAY BE USED TO IDENTIFY THERAPEUTICALLY ACTIONABLE ALTERATIONS³

ASCO recommends multigene panel-based genomic testing or NGS for:

Patients eligible for an approved genomic	Patients potentially eligible for more than 1
biomarker-linked therapy	approved genomic biomarker–linked therapy
To detect tumor-agnostic actionable biomarkers like dMMR and/or MSI-H, TMB-H, and <i>NTRK</i> fusions, which may not be detected by single-gene tests	To provide the most efficient use of limited tumor biopsy tissue

ASCO recommends using NGS for the most efficient utilization of limited biopsy tissue; it may allow simultaneous testing for multiple approved targeted therapies

dMMR, deficient mismatch repair; FISH, fluorescence in situ hybridization; IHC, immunohistochemistry; MSI-H, microsatellite instability-high; NTRK, neurotrophic tyrosine receptor kinase; RT-PCR, real-time polymerase chain reaction; TMB-H, tumor mutation burden-high.





BIOMARKERS FOR TARGETED THERAPIES ARE DIFFERENT THAN BIOMARKERS FOR ICIS

Targeted Therapy Biomarkers





Images adapted with permission from Camidge DR et al. Nat Rev Clin Oncol. 2019;16(6):341-355.

Targeted therapies inhibit cells harboring a specific genomic alteration or protein³

Responses to targeted therapies may be primarily influenced by the presence of a driver alteration assumed to be present in most tumor cells⁶⁷⁻⁷⁰

Biomarkers for targeted therapies⁶⁷:



May be categorial or continuous depending on the alteration (eg, mutation or amplification)^{68,71,a}



Are assumed to be present in most tumor cells⁶⁸

*Except for gene amplifications, which are continuous. mRNA, messenger RNA.



BIOMARKERS FOR TARGETED THERAPIES ARE DIFFERENT THAN BIOMARKERS FOR ICIS ICI Biomarkers



ICIs reduce T-cell exhaustion by disrupting the immune checkpoint⁷²⁻⁷⁵

Responses to ICIs may be influenced by complex interactions between multiple different cell types^{67,76}

Biomarkers for ICIs are⁶⁷:



Continuous with arbitrary cutoffs⁷⁷⁻⁷⁹



Spatially and temporally variable^{79,80}

PD-1, programmed cell death protein 1; PD-L1, programmed death ligand 1.

OVERVIEW OF SELECT BIOMARKER TESTING CHALLENGES



^aBased on a meta-analysis of 61 studies including more than 5,700 patients with metastatic colorectal cancer.³² ^bBased on the 2017 National Survey of Precision Medicine in Cancer Treatment by the National Cancer Institute. A total of 1,281 medical oncologists participated in this survey.³²



OVERVIEW OF SELECT BIOMARKER TESTING CHALLENGES

Extensive TAT

In some cancers with multiple biomarkers, studies suggest sequential single-gene testing may contribute to tissue exhaustion, potentially leading to:



Patients *not* receiving testing for all biomarkers



Prolonged TAT for all biomarkers (relative to a multigene panel)^{87,88}

Multigene panels may have TATs of

>10 days^{**}

Intended to depict biomarker testing methodologies. When testing for therapy selection, please consult product prescribing information and FDA approved CDx.



POTENTIAL SOLUTIONS TO OVERCOME CHALLENGES

A

Failure to obtain sufficient tissue during biopsy

ROSE May Improve Biopsy Yield

Implementation of ROSE has been associated with an **increase in diagnostic yield**



 With ROSE at a single center, presence of tumor material was confirmed in 86% of biopsies, 96% of which were sufficient for molecular testing^{95,c} in cytology
 procedures^{93,a}

Up to

in endobronchial ultrasound– guided transbronchial needle aspiration (EBUS-TBNA) procedures^{94,b}

- In one study with ROSE, ≈98% of samples obtained were deemed adequate^{96,d}
 - Some studies report that the diagnostic yield and accuracy were comparable in procedures done with and without ROSE^{97,98}

ROSE Stains Show Different Cytologic Details





Extensive **TAT**

Reflex Testing Can Help Streamline Biomarker Testing Reflex testing is the automatic addition of tests in the SOPs by pathologists⁹⁹ in specific situations, such as:

 $^{\sigma}$ An equivocal HER2 IHC result in breast cancer45

 $\left(\begin{smallmatrix} \\ 0 \\ 0 \\ 0 \\ \end{array}\right)$

Reflex testing may be integrated into the electronic health record¹⁰⁰ Reflex testing is dependent on the cancer type, staging, and institution protocol^{101,102}

Reflex Testing May Reduce TAT^{102,103}

A retrospective review of 166 patients diagnosed with lung adenocarcinoma between 2016 and 2018 at a community center assessed biomarker testing rates and TATs for molecular testing¹⁰³

Reflex ordered testing was implemented in February 2017¹⁰³

TATs were compared before and after reflex testing implementation¹⁰³

TAT was defined as the date of the anatomic pathology report confirming lung adenocarcinoma diagnosis to the date of the final molecular diagnostics report¹⁰³

TAT before reflex testing¹⁰³

TAT with reflex testing¹⁰³

15.6 days

Reduced TAT after reflex testing has been observed in other tumor types as well¹⁰²

HER2, human epidermal growth factor receptor 2; SOPs, standard operating procedures.

5**2.6** days



POTENTIAL SOLUTIONS TO OVERCOME CHALLENGES (CONTINUED)



Overall complexities associated with biomarker testing

MTBs May Help Navigate the Complexities of Precision Oncology

Cancer treatment recommendations from MTBs may be based on many factors, including¹⁰⁴:
Tumor type

Molecular alterations
Performance status
Comorbidities

Many specialties may be part of an MTB to help foster discussion¹⁰⁴
Image: Ima

• Pharmacists

Real-world evidence from a retrospective review of 782 patients with solid tumors tested with NGS in a tertiary care center suggests MTBs may help in appropriate and actionable clinical decision-making¹⁰⁵

MTB, molecular tumor board.



REFERENCES

1. Falzone L et al. Front Pharmacol. 2018;9:1300. 2. Dobosz P, Dzieciątkowski T. Front Immunol. 2019;10:2965. 3. Chakravarty D et al. J Clin Oncol. 2022;40(11):1231-1258. **4.** Sawyers CL. *Cell.* 2019;179(1):8-12. **5.** Vamathevan J et al. Nat Rev Drug Discov. 2019;18(6):463-477. **6.** Yu D et al. *Mol Cancer.* 2022;21(1):56. 7. Wang Y et al. Front Oncol. 2021;11:683419. 8. Hung K et al. J Exp Med. 1998;188(12):2357-2368. 9. Dudley ME et al. Science. 2002;298(5594):850-854. 10. Lassen UN et al. Future Oncol. 2021;17(30):3995-4009. 11. Hess LM et al. JTO Clin Res Rep. 2022;3(6):100336. 12. National Cancer Institute. TCGA timeline and milestones. https://www.cancer.gov/ about-nci/organization/ccg/research/structural-genomics/tcga/history/ timeline. Accessed April 13, 2022. 13. National Cancer Institute. The Cancer Genome Atlas project. https://www.cancer.gov/about-nci/organization/ccg/ research/structural-genomics/tcga. Accessed April 13, 2022. 14. Bailey MH et al. Cell. 2018;173(2):371-385.e18. 15. National Human Genome Research Institute. The Human Genome Project. https://www.genome.gov/humangenome-project. Accessed April 13, 2022. 16. National Human Genome Research Institute. The cost of sequencing a human genome. https://www. genome.gov/about-genomics/fact-sheets/Sequencing-Human-Genome-cost. Accessed April 1, 2022. **17.** Zhong Y et al. Ann Lab Med. 2021;41(1):25-43. **18.** Vogel CL et al. J Clin Oncol. 2002;20(3):719-726. **19.** US Food and Drug Administration. List of cleared or approved companion diagnostic devices (in vitro and imaging tools). https://www.fda.gov/medical-devices/in-vitrodiagnostics/list-cleared-or-approved-companion-diagnostic-devices-in-vitroand-imaging-tools. Accessed June 9, 2022. 20. Hodi FS et al. N Engl J Med. 2010;363(8):711-723. 21. Marcus L et al. Clin Cancer Res. 2019;25(13):3753-3758. 22. Seligson ND et al. Clin Pharmacol Ther. 2021;109(2):334-342. 23. Sochacka-Ćwikła A et al. *Molecules*. 2022;27(7):2259. 24. Drilon A et al. *N Engl J Med*. 2018;378(8):731-739. 25. Olivier T et al. *JAMA Netw* Open. 2021;4(12):e2138793. 26. OncoKB. Actionable genes. https://www. oncokb.org/actionableGenes#levels=1§ions=Tx. Accessed June 13, 2022. 27. De Las Casas LE, Hicks DG. Am J Clin Pathol. 2021;155(6):781-792. **28.** Saini KS et al. Ann Oncol. 2012;23(4):853-859. **29.** Danesi R et al. ESMO Open. 2021;6(2):100040. 30. Ersek JL et al. Am Soc Clin Oncl Educ Book. 2018;38:188-196. **31**. Cree IA et al. J Clin Pathol. 2014;67(11):923-931. **32**. Roberts MC et al. JCO Precis Oncol. 2021;5:PO.20.00431. doi: . **33**. Association of Community Cancer Centers. Precision medicine: integration of pathology with the cancer care team. https://www.accc-cancer.org/ docs/projects/landscape-of-pathology/pathology/patient-centered-care. pdf?sfvrsn=8173e1cb_2. Accessed March 21, 2022. **34**. Mikubo M et al. *J Thorac Oncol.* 2020;15(1):130-137. **35**. Frampton GM et al. *Nat Biotechnol.* 2013;31(11):1023-1031. **36**. Smits AJJ et al. *Mod Pathol.* 2014;27(2):168-174. 37. Stefanovic S et al. Oncotarget. 2017;8(31):51416-51428. 38. Bhullar DS et al. EBioMedicine. 2019;40:363-374. 39. Bardelli A et al. Proc Natl Acad Sci U S A. 2001;98(10):5770-5775. 40. Sidransky D et al. Nature. 1992;355(6363):846-847. 41. Yachida S et al. Nature. 2010;467(7319):1114-1117. 42. Singh VM et al. Ann Diagn Pathol. 2013;17(4):322-326. 43. Referenced with permission from the NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines®) for Breast Cancer V.4.2022. © National Comprehensive Cancer Network, Inc. 2022. All rights reserved. Accessed June 21, 2022. To view the most recent and complete version of the guideline, go online to NCCN.org. NCCN makes no warranties of any kind whatsoever regarding their content, use or application and disclaims any responsibility for their application or use in any way. 44. Shachar SS et al. *Clin Breast Cancer*. 2016;16(3):e43-e48.45. Swanton C. *Cancer Res*. 2012;72(19):4875-4882. 46. Gerlinger M et al. Intratumor heterogeneity and branched evolution revealed by multiregion sequencing. N Engl J Med. 2012;366(10):883-892. 47. Rolfo C et al. J Thorac Oncol. 2021;16(10):1647-1662. **48.** Merker JD et al. J Clin Oncol. 2018;36(16):1631-1641. **49.** Tay TKY, Tan PH. Arch Pathol Lab Med. 2021;145(6):678-686. 50. Yeung C et al. Cancer Metastasis Rev. 2016;35(3):427-437. 51. Arneth B. BMC Cancer. 2018;18(1):527. 52. Iuliani M et al. Front Oncol. 2020;10:789. 53. Gerratana L et al. Curr Treat Options Oncol. 2019;20(8):68. **54.** Davis AA et al. EBioMedicine. 2020;58:102914. 55. Gerratana L et al. Eur J Cancer. 2021;143:147-157. 56. Bhadresha KP et al. J Bone Oncol. 2021;29:100374. 57. Hayes DF. J Clin Oncol. 2021;39(3):238-248. 58. Vidwans SJ et al. Oncoscience. 2014;1(10):614-623. 59. Pepe MS et al. J Natl Cancer Inst. 2001;93(14):1054-1061. 60. Jennings LJ et al. J Mol Diagn. 2017;19(3):341-365. 61. European Society for Medical Oncology. Multigene sequencing in breast cancer: ESMO biomarker factsheet. https:// oncologypro.esmo.org/education-library/factsheets-on-biomarkers/multigenesequencing-in-breast-cancer#page. Accessed April 26, 2022. 62. Veljovic M et al. ASCO 2015. Abstract e17698. 63. Lim C et al. Ann Oncol. 2015;26(7):1415-1421. 64. Hayes DF. Mol Oncol. 2015;9(5):960-966. 65. Parkinson DR et al. Clin Cancer Res. 2014;20(6):1428-1444. 66. US Food and Drug Administration. Developing and labeling in vitro companion diagnostic devices for a specific group of oncology therapeutic products: guidance for industry. https://www.fda.gov/regulatory-information/search-fda-guidancedocuments/developing-and-labeling-in-vitro-companion-diagnostic-devices-specific-group-oncology-therapeutic. Accessed May 6, 2022. **67**. Camidge DR

et al. Nat Rev Clin Oncol. 2019;16(6):341-355. **68**. de Bruin EC et al. Science. 2014;346(6206):251-256. **69**. Zhu CQ et al. J Clin Oncol. 2008;26(26):4268-4275. **70**. Cappuzzo F et al. J Natl Cancer Inst. 2005;97(9):643-655. **71**. Noonan SA et al. J Thorac Oncol. 2016;11(8):1293-1304. **72**. Grywalska E et al. Onco Targets Ther. 2018;11:6505-6524.

73. Tumeh PC et al. Nature. 2014;515(7528):568-571. 74. Forde PM et al. N Engl J Med. 2018;378(21):1976-1986. **75**. Hamid O et al. N Engl J Med. 2013;369:134-144. **76.** Qin A et al. Sci Rep. 2022;12(1):9054. **77.** Mok TSK et al. Lancet. 2019;393(10183):1819-1830. 78. Brahmer J et al. N Engl J Med. 2015;373(2):123-135. 79. Lantuejoul S et al. J Thorac Oncol. 2020;15(4):499-519. 80. Büttner R et al. J Clin Oncol. 2017;35(34):3867-3876. 81. Bhamidipati D et al. NPJ Precis Oncol. 2021;5(1):94. 82. Bhullar DS et al. EBIoMedicine. 2019;40:363-374. 83. Xing D et al. Hum Pathol. 2019;92:67-80. 84. Kolberg-Liedtke C et al. Breast Care (Basel). 2021;16(5):475-483. 85. Ptotkin E et al. Integration of pathology within the multidisciplinary cancer care team. ASCO 2019. Abstract 49. 86. Committee on Policy Issues in the Clinical Development and Use of Biomarkers for Molecularly Targeted Therapies; Board on Health Care Services; Institute of Medicine; National Academies of Sciences, Engineering, and Medicine. Biomarker Tests for Molecularly Targeted Therapies: Key to Unlocking Precision Medicine. Graig LA, Phillips JK, Moses HL, eds. Washington, DC: National Academies; 2016 87. Yu TM et al. Clin Lung Cancer. 2019;20(1):20-29.e8 88. Pennell NA et al. JCO Precis Oncol. 2019;3:PO.18.00356. 89. College of American Pathologists. Definition of synoptic reporting. https://documents.cap.org/documents/ Synoptic reporting interpartation of the synoptic reporting definition examples v4.0.pdf. Accessed March 18, 2022.
 Valenstein PN. Arch Pathol Lab Med. 2008;132(1):84-94. Elenitoba-Johnson KS. Annu Rev Pathol. 2020;15:97-121. **92**. Nussinov R et al. PLoS Comput Biol. 2019;15(3):e1006658. **93**. Azabdaftari G et al. Acta Cytol. 2010;54(2):132-137. 94. Gianella P et al. Acta Cytol. 2018;62(5-6):380-385. 95. Manzo JL et al. Cancer Cytopathol. 2018;126(7):481-489. 96. Casal RF et al. BMC Endocr Disord. 2014;14:88. 97. Lee LS et al. Dig Endosc. 2016;28(4):469-475. 98. Kappelle WFW et al. Am J Gastroenterol. 2018;113(5):677-685. 99. Murphy MJ. Ann Clin Biochem. 2021;58(2):75-77. 100. Lau-Min KS et al. JCO Precis Oncol. 2021;5:PO.20.00418. 101. Association of Community Cancer Centers. Understanding the integration of pathology with the cancer care team: survey highlights. https://www.accc-cancer.org/docs/projects/landscape-of-pathology/pathologyinfographicsup-final-online.pdf. Accessed April 26, 2022. D2. Seidman AD et al. Popul Health Manag. 2017;20(4):252-254. 103. Anand K et al. Clin Lung Cancer. 2020;21(5):437-442. 104. Luchini C et al. Trends Cancer. 2020;6(9):738-744. 105. Sadaps M et al. The impact of clinical decision making in a molecular tumor board at a tertiary care center. ASCO 2021. Abstract 3128.

SUMMARY



Precision oncology has contributed to improved care for patients³

Each member of the MDT is important to biomarker testing collaboration and communication²⁷



Improving MDT communication and collaboration may increase the number of patients receiving biomarker-informed care²⁷



Do you have the Knowledge Check that goes with this chapter?

Are you interested in

learning more about

Precision Medicine?

VISIT OUR WEBSITE!



■ <a>

 You'll find knowledge checks,

 additional resources, a digital version of this and other chapters, and more

www.hcp.novartis.com/precision-medicine



Looking to speak to a Precision Medicine Liaison? Scan this QR code

www.hcp.novartis.com/precision-medicine/contact-us



Novartis Pharmaceuticals Corporation East Hanover, New Jersey 07936-1080

