

Overview

Useful For

Predicting potential for toxicity to thiopurine drugs (6-mercaptopurine, 6-thioguanine, and azathioprine)

Testing Algorithm

For more information see:

- [Ulcerative Colitis and Crohn Disease Therapeutic Drug Monitoring Algorithm](#)
- [TPMT Testing in the Treatment of Inflammatory Bowel Disease Algorithm](#)

Special Instructions

- [Informed Consent for Genetic Testing](#)
- [Pharmacogenomic Association Tables](#)
- [Multiple Genotype Test List](#)
- [Informed Consent for Genetic Testing \(Spanish\)](#)
- [Ulcerative Colitis and Crohn Disease Therapeutic Drug Monitoring Algorithm](#)
- [TPMT Testing in the Treatment of Inflammatory Bowel Disease Algorithm](#)

Highlights

This test includes genotyping of *TPMT* and *NUDT15*, both of which affect metabolism of thiopurine drugs.

Method Name

Real-Time Polymerase Chain Reaction (PCR) With Allelic Discrimination Analysis

NY State Available

Yes

Specimen

Specimen Type

Varies

Ordering Guidance

For thiopurine methyltransferase (TPMT) enzyme activity testing, order TPMT3 / Thiopurine Methyltransferase Activity Profile, Erythrocytes; however, this test should also be ordered because TPMT enzyme activity testing cannot detect variants in *NUDT15*, which also impact thiopurine metabolism.

Specimen Required

Submit only 1 of the following specimens:

Patient Preparation: A previous hematopoietic stem cell transplant from an allogenic donor will interfere with testing.

Call 800-533-1710 for instructions for testing patients who have received a hematopoietic stem cell transplant.

Specimen Type: Whole blood

Container/Tube: Lavender top (EDTA)

Specimen Volume: 3 mL

Collection Instructions:

1. Invert several times to mix blood.
2. Send whole blood specimen in original tube. **Do not aliquot.**

Specimen Stability Information: Ambient (preferred) 4 days/Refrigerated 4 days/Frozen 4 days

Additional Information:

1. Specimens are preferred to be received within 4 days of collection. Extraction will be attempted for specimens received after 4 days, and DNA yield will be evaluated to determine if testing may proceed.
2. To ensure minimum volume and concentration of DNA is met, the requested volume must be submitted. Testing may be canceled if DNA requirements are inadequate.

Specimen Type: Cord blood

Container/Tube: Lavender top (EDTA)

Specimen Volume: 3 mL

Collection Instructions:

1. Invert several times to mix blood.
2. Send specimen in original tube. **Do not aliquot.**

Specimen Stability Information: Ambient (preferred) 4 days/Refrigerated 4 days/Frozen 4 days

Additional Information:

1. Specimens are preferred to be received within 4 days of collection. Extraction will be attempted for specimens received after 4 days, and DNA yield will be evaluated to determine if testing may proceed.
2. To ensure minimum volume and concentration of DNA is met, the requested volume must be submitted. Testing may be canceled if DNA requirements are inadequate.
3. While a properly collected cord blood sample may not be at risk for maternal cell contamination, unanticipated complications may occur during collection. Therefore, maternal cell contamination studies are recommended to ensure the test results reflect that of the patient tested and are available at an additional charge.
4. Order MATCC / Maternal Cell Contamination, Molecular Analysis, Varies on the maternal specimen.

Specimen Type: Saliva

Patient Preparation: Patient **should not** eat, drink, smoke, or chew gum 30 minutes prior to collection.

Supplies: Saliva Swab Collection Kit (T786)

Specimen Volume: 1 Swab

Collection Instructions: Collect and send specimen per kit instructions.

Specimen Stability Information: Ambient (preferred) 30 days/Refrigerated 30 days

Additional Information: Saliva specimens are acceptable but not recommended. Due to lower quantity/quality of DNA yielded from saliva, some aspects of the test may not perform as well as DNA extracted from a whole blood sample. When applicable, specific gene regions that were unable to be interrogated will be noted in the report. Alternatively, additional specimen may be required to complete testing.

Specimen Type: Extracted DNA

Container/Tube:

Preferred: Screw Cap Micro Tube, 2 mL with skirted conical base

Acceptable: Matrix tube, 1 mL

Collection Instructions:

- 1. The preferred volume is at least 100 mL at a concentration of 75 ng/mL.
- 2. Include concentration and volume on tube.

Specimen Stability Information: Frozen (preferred) 1 year/Ambient/Refrigerated

Additional Information: DNA must be extracted in a CLIA-certified laboratory or equivalent and must be extracted from a specimen type listed as acceptable for this test (including applicable anticoagulants). Our laboratory has experience with Chemagic, Puregene, Autopure, MagnaPure, and EZ1 extraction platforms and cannot guarantee that all extraction methods are compatible with this test. If testing fails, one repeat will be attempted, and if unsuccessful, the test will be reported as failed and a charge will be applied. If applicable, specific gene regions that were unable to be interrogated due to DNA quality will be noted in the report.

Forms

- 1. **New York Clients-Informed consent is required.** Document on the request form or electronic order that a copy is on file. The following documents are available:
 - [Informed Consent for Genetic Testing](#) (T576)
 - [Informed Consent for Genetic Testing-Spanish](#) (T826)
- 2. If not ordering electronically, complete, print, and send 1 of the following forms with the specimen:
 - [Neurology Specialty Testing Client Test Request](#) (T732)
 - [Gastroenterology and Hepatology Test Request](#) (T728)
 - [Therapeutics Test Request](#) (T831)

Specimen Minimum Volume

See Specimen Required

Reject Due To

All specimens will be evaluated at Mayo Clinic Laboratories for test suitability.

Specimen Stability Information

Specimen Type	Temperature	Time	Special Container
Varies	Varies		

Clinical & Interpretive

Clinical Information

The thiopurine drugs are purine antimetabolites that are useful in the treatment of acute lymphoblastic leukemia, autoimmune disorders (eg, Crohn disease, rheumatoid arthritis), and organ transplant recipients. The thiopurine drugs, 6-mercaptopurine, 6-thioguanine, and azathioprine are prodrugs that require intracellular activation to 6-thioguanine nucleotides (6-TGN). This activation is catalyzed by multiple enzymes. The cytotoxic effects of thiopurine drugs are achieved mainly through incorporation of 6-TGN into DNA and RNA. The pathway that leads to synthesis of active cytotoxic 6-TGN is in competition with inactivation pathways catalyzed by thiopurine methyltransferase (TPMT).

Evaluation of this pathway is important because the level of 6-TGN measured in red blood cells have been correlated with both thiopurine therapeutic efficacy and toxicity such as myelosuppression.

TPMT activity is inherited as a monogenic codominant trait, and variable TPMT activity is associated with *TPMT* genetic variants. The distribution of TPMT activity in red blood cells is trimodal in the population of people with European ancestry, with approximately 0.3% having deficient (undetectable) TPMT activity, 11% low (intermediate) activity, and 89% normal TPMT activity. The allele for normal TPMT activity (wild type) has been designated *TPMT**1. Four *TPMT* alleles, comprised of a combination of 3 different single-nucleotide variants, account for the majority of inactivating alleles in some ancestral populations, including Europeans: *TPMT**2, *TPMT**3A, and *TPMT**3C. Less frequently occurring *TPMT* alleles including *TPMT**4, *TPMT**5, *TPMT**8, and *TPMT**12 also have been implicated as deficiency alleles. If no *TPMT* variant alleles are detected by this assay, the most likely genotype is that of *TPMT**1/*1, although the presence of other rarer alleles cannot be excluded.

Nudix hydrolase (NUDT15) is thought to dephosphorylate the active metabolites of thiopurines, TGTP, and TdGTP, which prevents their incorporation into DNA and decreases their cytotoxic effects. Genetic variants in *NUDT15* that decrease this activity are strongly associated with thiopurine-related myelosuppression. NUDT15 deficiency is most common among East Asian (22.6%), South Asian (13.6%), and Native American populations (12.5%-21.2%). Studies in other populations are ongoing. This test evaluates variants associated with *NUDT15**2, *NUDT15**3, *NUDT15**4, and *NUDT15**5. If no *NUDT15* variant alleles are detected by this assay, the most likely genotype is that of *NUDT15**1/*1, although the presence of other rarer alleles cannot be excluded. Individuals with variants in both *TPMT* and *NUDT15* have been identified and were significantly more sensitive to mercaptopurine than individuals heterozygous for a variant in only one gene. Integration of both *TPMT* and *NUDT15* testing may allow for more accurate prediction of thiopurine-related toxicity risk to guide dosing, particularly among patients from diverse populations.

Table 1. TPMT Enzyme Activity of Individual Star Alleles

<i>TPMT</i> allele	cDNA nucleotide change (NM_000367.4)	Amino acid change	Effect on enzyme metabolism
*1	None (wild type)	None (wild type)	Normal function
*2	c.238G>C	p.Ala80Pro (p.A80P)	No activity
*3A	c.460G>A and c.719A>G	p.Ala154Thr (p.A154T) and p.Tyr240Cys (p.Y240C)	No activity
*3B	c.460G>A	p.Ala154Thr (p.A154T)	No activity
*3C	c.719A>G	p.Tyr240Cys (p.Y240C)	No activity
*4	c.626-1G>A	Not applicable, splice site	No activity
*5	c.146T>C	p.Leu49Ser (p.L49S)	No activity
*8	c.644G>A	p.Arg215His (p.R215H)	Reduced activity
*12	c.374C>T	p.Ser125Leu (p.S125L)	Reduced activity

Table 2. NUDT15 Enzyme Activity of Individual Star Alleles

<i>NUDT15</i> allele	cDNA nucleotide change (NM_018283.3)	Amino acid change	Effect on enzyme metabolism
*1	None (wild type)	None (wild type)	Normal activity

*2 or *3	c.415C>T	p.Arg139Cys (p.R139C)	No activity
*4	c.416G>A	p.Arg139His (p.R139H)	No activity
*5	c.52G>A	p.Val18Ile (p.V18I)	No activity

The US Food and Drug Administration, the Clinical Pharmacogenetics Implementation Consortium, and some professional societies recommend consideration of *TPMT* and *NUDT15* genotype testing or *TPMT* enzyme activity testing along with *NUDT15* genotype testing prior to the initiation of therapy with thiopurine drugs. There is substantial evidence linking *TPMT* and *NUDT15* genotypes to phenotypic variability. Dose adjustments based upon *TPMT* and *NUDT15* genotypes have reduced thiopurine-induced adverse effects without compromising desired antitumor and immunosuppressive therapeutic effects in several clinical settings.

Genotyping is not impacted by other medications known to inhibit *TPMT* activity. Complementary clinical testing is available to measure *TPMT* enzymatic activity in erythrocytes (*TPMT3* / Thiopurine Methyltransferase Activity Profile, Erythrocytes) if the clinician wants to check for lower *TPMT* enzyme activity, regardless of cause. Testing for *TPMT* enzyme activity is not impacted by variants in *NUDT15*.

Reference Values

An interpretive report will be provided.

Interpretation

The *TPMT* genotype, with associated star alleles, is assigned using standard allelic nomenclature as published by the *TPMT* Nomenclature Committee.(1) *NUDT15* genotype and associated star alleles are as described by Moriyama et al.(2) and catalogued in the Pharmacogene Variation Consortium (www.pharmvar.org).

For additional information regarding pharmacogenomic genes and their associated drugs, see the [Pharmacogenomics Associations Tables](#). This resource also includes information regarding enzyme inhibitors and inducers, as well as potential alternate drug choices.

Cautions

Rare variants may be present that could lead to false-negative or false-positive results. If no *TPMT* variant alleles are detected by this assay, the most likely genotype is that of *TPMT**1/*1 although the presence of other rarer alleles cannot be excluded. In addition, if no *NUDT15* variant alleles are detected by this assay, the most likely genotype is that of *NUDT15**1/*1, although the presence of other rarer alleles cannot be excluded.

If genotype results obtained do not match the clinical findings, additional testing should be considered for thiopurine methyltransferase enzyme activity (*TPMT3* / Thiopurine Methyltransferase Activity Profile, Erythrocytes). A corresponding activity assay for *NUDT15* is not currently available.

Specimens may contain donor DNA if obtained from patients who received non-leukoreduced blood transfusions or allogeneic hematopoietic stem cell transplantation. Results from specimens obtained under these circumstances may not accurately reflect the recipient's genotype. For individuals who have received blood transfusions, the genotype usually reverts to that of the recipient within 6 weeks. For individuals who have received allogeneic hematopoietic stem cell transplantation, a pretransplant DNA specimen is recommended for testing.

The results do not rule out the possibility that a patient harbors another variant in *TPMT*, *NUDT15*, or another gene that

can impact drug response or side effects. These genotyping procedures will not distinguish between *TPMT* *1/*3A from the rare *TPMT* *3B/*3C genotype, which has an estimated frequency of 1:120,890. This rare genotype is associated with low enzyme activity. Enzyme activity evaluation is necessary to definitively identify this rare genotype (TPMT3 / Thiopurine Methyltransferase Activity Profile, Erythrocytes).

This test will not detect all *TPMT* or *NUDT15* genetic variants. A negative result does not rule out the possibility of toxicity if thiopurines are used, since multiple factors (eg, other genetic factors, drug-drug interactions) are known to play a role. Co-prescription of allopurinol might inhibit TPMT activity. Other drugs that have been shown to inhibit TPMT activity include naproxen, ibuprofen, ketoprofen, furosemide, sulfasalazine, mesalamine, olsalazine, mefenamic acid, thiazide diuretics, and benzoic acid inhibitors.

Clinical Reference

1. TPMT nomenclature committee (TPMT Alleles): Table of TPMT Alleles. Linköping University; Updated November 2022. Accessed March 19, 2025. Available at <https://liu.se/en/research/tpmt-nomenclature-committee>
2. Moriyama T, Nishii R, Perez-Andreu V, et al. NUDT15 polymorphisms alter thiopurine metabolism and hematopoietic toxicity. *Nat Genet*. 2016;48(4):367-373. doi:10.1038/ng.3508
3. Appell ML, Berg J, Duley J, et al. Nomenclature for alleles of the thiopurine methyltransferase gene. *Pharmacogenet Genomics*. 2013;23(4):242-248. doi:10.1097/FPC.0b013e32835f1cc0
4. Nguyen CM, Mendes MA, Ma JD. Thiopurine methyltransferase (TPMT) genotyping to predict myelosuppression risk. *PLoS Curr*. 2011;3:RRN1236. doi:10.1371/currents.RRN1236
5. Relling MV, Schwab M, Whirl-Carrillo M, et al. Clinical Pharmacogenetics Implementation Consortium guideline for thiopurine dosing based on TPMT and NUDT15 genotypes: 2018 Update. *Clin Pharmacol Ther*. 2019;105(5):1095-1105. doi:10.1002/cpt.1304
6. Weinshilboum R. Thiopurine pharmacogenetics: clinical and molecular studies of thiopurine methyltransferase. *Drug Metab Dispos*. 2001;29(4 Pt 2):601-605
7. Zaza G, Cheok M, Krynetskaia N, et al. Thiopurine pathway. *Pharmacogenet Genomics*. 2010;20(9):573-574. doi:10.1097/FPC.0b013e328334338f
8. Sterner RM, Hall PL, Matern D, Black JL, Moyer AM. Genotype and Phenotype Correlation of the TPMT*8 Allele in Thiopurine Metabolism. *J Mol Diagn*. 2024;26(11):988-994. doi:10.1016/j.jmoldx.2024.07.005
9. Pratt VM, Cavallari LH, Fulmer ML, et al. TPMT and NUDT15 Genotyping Recommendations: A Joint Consensus Recommendation of the Association for Molecular Pathology, Clinical Pharmacogenetics Implementation Consortium, College of American Pathologists, Dutch Pharmacogenetics Working Group of the Royal Dutch Pharmacists Association, European Society for Pharmacogenomics and Personalized Therapy, and Pharmacogenomics Knowledgebase. *J Mol Diagn*. 2022;24(10):1051-1063. doi:10.1016/j.jmoldx.2022.06.007

Performance

Method Description

Genomic DNA is extracted from whole blood or saliva. Genotyping for the *TPMT* and *NUDT15* alleles is performed using a polymerase chain reaction (PCR)-based 5'-nuclease assay. Fluorescently labeled detection probes anneal to the target DNA. PCR is used to amplify the segment of DNA that contains the polymorphism. If the detection probe is an exact match to the target DNA, the 5'-nuclease polymerase degrades the probe, the reporter dye is released from the effects

of the quencher dye, and a fluorescent signal is detected. Genotypes are assigned based on the allele-specific fluorescent signals that are detected.(Unpublished Mayo method)

PDF Report

No

Day(s) Performed

Monday through Friday

Report Available

2 to 4 days

Specimen Retention Time

Whole blood/Saliva: 30 days (if available); Extracted DNA: 3 months

Performing Laboratory Location

Rochester

Fees & Codes

Fees

- Authorized users can sign in to [Test Prices](#) for detailed fee information.
- Clients without access to Test Prices can contact [Customer Service](#) 24 hours a day, seven days a week.
- Prospective clients should contact their account representative. For assistance, contact [Customer Service](#).

Test Classification

This test was developed and its performance characteristics determined by Mayo Clinic in a manner consistent with CLIA requirements. It has not been cleared or approved by the US Food and Drug Administration.

CPT Code Information

0034U

LOINC® Information

Test ID	Test Order Name	Order LOINC® Value
TPNUQ	TPMT and NUDT15 Genotype, V	93193-1

Result ID	Test Result Name	Result LOINC® Value
610159	TPMT Genotype	41048-0
610160	TPMT Phenotype	79713-4
610161	NUDT15 Genotype	93194-9
610162	NUDT15 Phenotype	93195-6
610163	Interpretation	69047-9

Test Definition: TPNUQ

Thiopurine Methyltransferase (TPMT) and
Nudix Hydrolase (NUDT15) Genotyping, Varies

610164	Additional Information	48767-8
610165	Method	85069-3
610166	Disclaimer	62364-5
610167	Reviewed by	18771-6