



FEP Medical Policy Manual

FEP 2.04.126 Germline Genetic Testing for Gene Variants Associated With Breast Cancer in Individuals at High Breast Cancer Risk (CHEK2, ATM, and BARD1)

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Related Policies:

- 2.04.02 - Germline Genetic Testing for Hereditary Breast/Ovarian Cancer Syndrome and Other High-Risk Cancers (BRCA1, BRCA2, PALB2)
- 2.04.08 - Genetic Testing for Lynch Syndrome and Other Inherited Colon Cancer Syndromes
- 2.04.101 - Genetic Testing for Li-Fraumeni Syndrome
- 2.04.149 - Molecular Testing for Germline Variants Associated with Ovarian Cancer (BRIP1, RAD51C, RAD51D, NBN)
- 2.04.88 - Genetic Testing for PTEN Hamartoma Tumor Syndrome
- 2.04.93 - Genetic Cancer Susceptibility Panels Using Next Generation Sequencing
- 6.01.29 - Magnetic Resonance Imaging for Detection and Diagnosis of Breast Cancer

Germline Genetic Testing for Gene Variants Associated With Breast Cancer in Individuals at High Breast Cancer Risk (CHEK2, ATM, and BARD1)

Description

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It is estimated that 3% to 5% of women presenting for assessment for hereditary breast/ovarian cancer risk have a variant in a gene that moderately increases the risk of cancer. *CHEK2*, *ATM*, and *BARD1* variants are considered to be of moderate penetrance. Female carriers of *CHEK2*, *ATM*, and *BARD1* have an approximately 2- to 4-fold increased risk of developing breast cancer compared with the general population. Risk estimates may be higher in patients with a family history of breast cancer or a family history of a specific variant.

OBJECTIVE

The objective of this review is to determine whether testing for *CHEK2*, *ATM*, and *BARD1* variants in individuals with a risk of hereditary breast/ovarian cancer improves the net health outcome.

POLICY STATEMENT

Testing for *CHEK2*, *ATM*, and *BARD1* variants in the assessment of breast cancer risk is considered **investigational**.

POLICY GUIDELINES

Criteria for Genetic Risk Evaluation

The National Comprehensive Cancer Network (NCCN) provides criteria for genetic risk evaluation for individuals with no history of breast cancer and for those with breast cancer. Updated versions of the criteria are available on the NCCN website.

The recommended testing strategy for *BRCA1*, *BRCA2*, and *PALB2* is described in review [2.04.02 \(genetic testing for hereditary breast/ovarian cancer syndrome\)](#).

Genetic Counseling

Genetic counseling is primarily aimed at patients who are at risk for inherited disorders, and experts recommend formal genetic counseling in most cases when genetic testing for an inherited condition is considered. The interpretation of the results of genetic tests and the understanding of risk factors can be very difficult and complex. Therefore, genetic counseling will assist individuals in understanding the possible benefits and harms of genetic testing, including the possible impact of the information on the individual's family. Genetic counseling may alter the utilization of genetic testing substantially and may reduce inappropriate testing. Genetic counseling should be performed by an individual with experience and expertise in genetic medicine and genetic testing methods.

BENEFIT APPLICATION

Experimental or investigational procedures, treatments, drugs, or devices are not covered (See General Exclusion Section of brochure).

Screening (other than the preventive services listed in the brochure) is not covered. Please see Section 6 General exclusions.

Benefits are available for specialized diagnostic genetic testing when it is medically necessary to diagnose and/or manage a patient's existing medical condition. Benefits are not provided for genetic panels when some or all of the tests included in the panel are not covered, are experimental or investigational, or are not medically necessary.

FDA REGULATORY STATUS

Clinical laboratories may develop and validate tests in-house and market them as a laboratory service; laboratory-developed tests must meet the general regulatory standards of the Clinical Laboratory Improvement Amendments. *CHEK2*, *ATM*, and *BARD1* testing are available under the auspices of the Clinical Laboratory Improvement Amendments. Laboratories offering to test and voluntarily listing is available through the National Center for Biotechnology Genetic Testing Registry. Laboratories that offer laboratory-developed tests must be licensed by the Clinical Laboratory Improvement Amendments for high-complexity testing. To date, the U.S. Food and Drug Administration has chosen not to require any regulatory review of this test.

Customized next-generation sequencing panels provide simultaneous analysis of multiple cancer predisposition genes, and typically include both moderate- and high-penetrant genes.

RATIONALE

Summary of Evidence

For individuals with risk of hereditary breast cancer/ovarian cancer (HBOC) who receive genetic testing for a checkpoint kinase 2 (*CHEK2*) variant, the evidence includes studies of variant prevalence and studies of breast cancer risk. Relevant outcomes are overall survival (OS), disease-specific survival, and test validity. The available studies on clinical validity have demonstrated that *CHEK2* variants are of moderate penetrance, and confer a risk of breast cancer 2 to 4 times that of the general population. Direct evidence for the clinical utility of genetic testing for *CHEK2* variants in individuals with risk of HBOC was not identified. It is unclear whether the relative risk (RR) associated with the moderate penetrance variants would increase risk enough beyond that already conferred by familial risk to change screening behavior. In contrast to high-penetrance variants, there is unlikely to be a similar benefit-to-risk calculus for risk-reducing mastectomy in women with a moderate penetrance variant such as *CHEK2*. The evidence is insufficient to determine that the technology results in an improvement in the net health outcome.

For individuals with risk of HBOC who receive genetic testing for an ataxia-telangiectasia mutated (*ATM*) variant, the evidence includes studies of variant prevalence and studies of breast cancer risk. Relevant outcomes are OS, disease-specific survival, and test validity. The available studies on clinical validity have demonstrated that *ATM* variants are of moderate penetrance; moreover, *ATM* variants confer a risk of breast cancer 2 to 4 times that of the general population. Direct evidence for the clinical utility of genetic testing for *ATM* variants in individuals with risk of HBOC was not identified. It is unclear whether the RR associated with the moderate penetrance variants would increase risk enough beyond that already conferred by familial risk to change screening behavior. In contrast to high-penetrance variants, there is unlikely to be a similar benefit-to-risk calculus for preventive interventions in women with a moderate penetrance variant such as *ATM*. The evidence is insufficient to determine that the technology results in an improvement in the net health outcome.

For individuals with risk of HBOC who receive genetic testing for a BRCA1-associated RING [Really Interesting New Gene] domain (*BARD1*) variant, the evidence includes studies of variant prevalence and studies of breast cancer risk. Relevant outcomes are OS, disease-specific survival, and test validity. The available studies on clinical validity have demonstrated that *BARD1* variants are of low to moderate penetrance; *BARD1* variants confer a risk of breast cancer about 2 to 3 times that of the general population. Direct evidence for the clinical utility of genetic testing for *BARD1* variants in individuals with a risk of HBOC was not identified. It is unclear whether the RR associated with the low to moderate penetrance variants would increase risk enough beyond that already conferred by familial risk to change screening behavior. In contrast to high-penetrance variants, there is unlikely to be a similar benefit-to-risk calculus for preventive interventions in women with a low to moderate penetrance variant such as *BARD1*. The evidence is insufficient to determine that the technology results in an improvement in the net health outcome.

SUPPLEMENTAL INFORMATION

Practice Guidelines and Position Statements

Guidelines or position statements will be considered for inclusion in 'Supplemental Information' if they were issued by, or jointly by, a US professional society, an international society with US representation, or National Institute for Health and Care Excellence (NICE). Priority will be given to guidelines that are informed by a systematic review, include strength of evidence ratings, and include a description of management of conflict of interest.

American College of Radiology

The American College of Radiology (ACR) has established Appropriateness Criteria for breast cancer screening (Table 1).⁴⁸ This includes high-risk women with a *BRCA* gene mutation and their untested first-degree relatives, women with a history of chest irradiation between 10 to 30 years of age, and women with 20% or greater lifetime risk of breast cancer as follows:

Table 1. American College of Radiology Appropriateness Criteria for Breast Cancer Screening in High-Risk Women

Screening Procedure	Appropriateness Category
Mammography	Usually appropriate
DBT	Usually appropriate
Breast MRI without and with IV contrast	May be appropriate

Breast US	May be appropriate
FDG-PEM	Usually not appropriate
Sestamibi MBI	Usually not appropriate
Breast MRI without IV contrast	Usually not appropriate

DBT: digital breast tomosynthesis; FDG-PEM: fludeoxyglucose positron emission mammography; IV: intravenous; MBI: molecular breast imaging; MRI: magnetic resonance imaging; US: ultrasound.

Specific recommendations for *CHEK2*, *ATM*, and *BARD1* variant carriers are not available.

American Society of Breast Surgeons

A consensus guideline on genetic testing for hereditary breast cancer was updated in February 2019.⁴⁹ Guidelines state that genetic testing should be made available to all individuals with a personal history of breast cancer and that such testing should include *BRCA1/BRCA2* and *PALB2*, with other genes as appropriate for the clinical scenario and patient family history. Furthermore, individuals who had previous genetic testing may benefit from updated testing. Finally, genetic testing should be made available to individuals without a personal history of breast cancer when they meet National Comprehensive Cancer Network (NCCN) guideline criteria. The guidelines also note that variants of uncertain significance are not clinically actionable.

For individuals with mutations in *ATM* and *CHEK2*, enhanced screening is recommended, however, the data are not sufficient to support risk-reducing mastectomy in the absence of other factors such as strong family history. For individuals with *BARD1* mutations, evidence is insufficient to support change in breast cancer risk management based on the presence of a mutation alone.

National Comprehensive Cancer Network

The NCCN (v.3.2023) guidelines on genetic/familial high-risk assessment for breast and ovarian cancer review single-gene tests for *CHEK2*, *ATM*, and *BARD1*.⁵⁰ The guidelines state that for those that meet hereditary cancer testing criteria, testing for a specific familial pathogenic/likely pathogenic variant may be recommended for appropriate genes. For individuals who meet criteria with no known familial variants, comprehensive testing of a multigene panel may be considered. This testing may consider a number of genes, including but not limited to *CHEK2*, *ATM*, and *BARD1*. However, the inclusion of certain genes in the guideline does not imply the endorsement "for or against multigene testing for moderate-penetrance genes" and there are limited data on the degree of cancer risk associated with some genes in multigene panels. Testing an affected family member first has the highest likelihood of a positive result. The guidelines state that the panel recommends an annual mammogram for women with *CHEK2*, *ATM*, or *BARD1* mutations beginning at age 40, with consideration of annual breast magnetic resonance imaging. The guidelines also state there is insufficient evidence to draw conclusions on risk-reducing mastectomy in individuals with *CHEK2*, *ATM*, or *BARD1* mutations and that patients should be managed based on family history.

The NCCN guidelines on breast cancer screening and diagnosis (v.1.2023)⁵¹, recommend the following:

- Annual mammogram.
- Annual breast magnetic resonance imaging if the patient has >20% risk of breast cancer based on models largely dependent on family history.
- Consideration of a risk-reducing strategies based on family history.

U.S. Preventive Services Task Force Recommendations

No U.S. Preventive Services Task Force recommendations for *CHEK2*, *ATM*, and *BARD1* variant testing have been identified.

Medicare National Coverage

There is no national coverage determination. In the absence of a national coverage determination, coverage decisions are left to the discretion of local Medicare carriers.

REFERENCES

1. National Cancer Institute, Surveillance Epidemiology and End Results Program. Cancer Stat Facts: Common Cancer Sites. 2023 <https://seer.cancer.gov/statfacts/html/common.html>. Accessed July 19, 2023.
2. National Cancer Institute, Surveillance Epidemiology and End Results Program. Cancer Stat Facts: Female Breast Cancer. n.d.; <https://seer.cancer.gov/statfacts/html/breast.html>. Accessed July 15, 2023.
3. National Cancer Institute. BRCA Mutations: Cancer Risk and Genetic Testing. November 19, 2020; <https://www.cancer.gov/about-cancer/causes-prevention/genetics/brca-fact-sheet>. Accessed July 18, 2023.
4. American Society of Clinical Oncology. Breast Cancer in Men: Risk Factors. 2021. <https://www.cancer.net/cancer-types/breast-cancer-men/risk-factors>. Accessed June 18, 2023.
5. Śniadecki M, Brzeziński M, Darecka K, et al. BARD1 and Breast Cancer: The Possibility of Creating Screening Tests and New Preventive and Therapeutic Pathways for Predisposed Women. *Genes (Basel)*. Oct 24 2020; 11(11). PMID 33114377
6. Alenezi WM, Fierheller CT, Recio N, et al. Literature Review of BARD1 as a Cancer Predisposing Gene with a Focus on Breast and Ovarian Cancers. *Genes (Basel)*. Jul 27 2020; 11(8). PMID 32726901
7. Suszynska M, Kluzniak W, Wokolorczyk D, et al. BARD1 is a Low/Moderate Breast Cancer Risk Gene: Evidence Based on An Association Study of the Central European p.Q564X Recurrent Mutation. *Cancers (Basel)*. May 28 2019; 11(6). PMID 31142030
8. Vysotskaia V, Kaseniit KE, Bucheit L, et al. Clinical utility of hereditary cancer panel testing: Impact of PALB2, ATM, CHEK2, NBN, BRIP1, RAD51C, and RAD51D results on patient management and adherence to provider recommendations. *Cancer*. Feb 01 2020; 126(3): 549-558. PMID 31682005
9. Apostolou P, Fostira F. Hereditary breast cancer: the era of new susceptibility genes. *Biomed Res Int*. 2013; 2013: 747318. PMID 23586058
10. Easton DF, Pharoah PD, Antoniou AC, et al. Gene-panel sequencing and the prediction of breast-cancer risk. *N Engl J Med*. Jun 04 2015; 372(23): 2243-57. PMID 26014596
11. Richards S, Aziz N, Bale S, et al. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology. *Genet Med*. May 2015; 17(5): 405-24. PMID 25741868
12. Kurian AW, Antoniou AC, Domchek SM. Refining Breast Cancer Risk Stratification: Additional Genes, Additional Information. *Am Soc Clin Oncol Educ Book*. 2016; 35: 44-56. PMID 27249685
13. Yadav S, LaDuca H, Polley EC, et al. Racial and Ethnic Differences in Multigene Hereditary Cancer Panel Test Results for Women With Breast Cancer. *J Natl Cancer Inst*. Oct 01 2021; 113(10): 1429-1433. PMID 33146377
14. Cybulski C, Wokolorczyk D, Jakubowska A, et al. Risk of breast cancer in women with a CHEK2 mutation with and without a family history of breast cancer. *J Clin Oncol*. Oct 01 2011; 29(28): 3747-52. PMID 21876083
15. Hu C, Hart SN, Gnanaolivu R, et al. A Population-Based Study of Genes Previously Implicated in Breast Cancer. *N Engl J Med*. Feb 04 2021; 384(5): 440-451. PMID 33471974
16. Schottenfeld D, Fraumeni JF. *Cancer epidemiology and prevention*. 3rd ed. New York: Oxford University Press; 2006.
17. Singletary SE. Rating the risk factors for breast cancer. *Ann Surg*. Apr 2003; 237(4): 474-82. PMID 12677142
18. Antoniou AC, Pharoah PP, Smith P, et al. The BOADICEA model of genetic susceptibility to breast and ovarian cancer. *Br J Cancer*. Oct 18 2004; 91(8): 1580-90. PMID 15381934
19. Berry DA, Iversen ES, Gudbjartsson DF, et al. BRCAPRO validation, sensitivity of genetic testing of BRCA1/BRCA2, and prevalence of other breast cancer susceptibility genes. *J Clin Oncol*. Jun 01 2002; 20(11): 2701-12. PMID 12039933
20. Nelson HD, Fu R, Goddard K, et al. Risk Assessment, Genetic Counseling, and Genetic Testing for BRCA- Related Cancer (AHRQ Publication No. 12-05164-EF-1). Rockville, MD: Agency for Healthcare Research and Quality; 2013.
21. Nelson HD, Pappas M, Zakher B, et al. Risk assessment, genetic counseling, and genetic testing for BRCA-related cancer in women: a systematic review to update the U.S. Preventive Services Task Force recommendation. *Ann Intern Med*. Feb 18 2014; 160(4): 255-66. PMID 24366442
22. Balmaa J, Digiovanni L, Gaddam P, et al. Conflicting Interpretation of Genetic Variants and Cancer Risk by Commercial Laboratories as Assessed by the Prospective Registry of Multiplex Testing. *J Clin Oncol*. Dec 2016; 34(34): 4071-4078. PMID 27621404
23. Suszynska M, Klonowska K, Jasinska AJ, et al. Large-scale meta-analysis of mutations identified in panels of breast/ovarian cancer-related genes - Providing evidence of cancer predisposition genes. *Gynecol Oncol*. May 2019; 153(2): 452-462. PMID 30733081
24. Yang Y, Zhang F, Wang Y, et al. CHEK2 1100delC variant and breast cancer risk in Caucasians: a meta-analysis based on 25 studies with 29,154 cases and 37,064 controls. *Asian Pac J Cancer Prev*. 2012; 13(7): 3501-5. PMID 22994785
25. Schmidt MK, Hogervorst F, van Hien R, et al. Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. *J Clin Oncol*. Aug 10 2016; 34(23): 2750-60. PMID 27269948
26. Weischer M, Bojesen SE, Ellervik C, et al. CHEK2*1100delC genotyping for clinical assessment of breast cancer risk: meta-analyses of 26,000 patient cases and 27,000 controls. *J Clin Oncol*. Feb 01 2008; 26(4): 542-8. PMID 18172190
27. Southey MC, Dowty JG, Riaz M, et al. Population-based estimates of breast cancer risk for carriers of pathogenic variants identified by gene-panel testing. *NPJ Breast Cancer*. Dec 09 2021; 7(1): 153. PMID 34887416
28. Li N, Lim BWX, Thompson ER, et al. Investigation of monogenic causes of familial breast cancer: data from the BEACCON case-control study. *NPJ Breast Cancer*. Jun 11 2021; 7(1): 76. PMID 34117267
29. Nguyen-Dumont T, Dowty JG, Steen JA, et al. Population-Based Estimates of the Age-Specific Cumulative Risk of Breast Cancer for Pathogenic Variants in CHEK2 : Findings from the Australian Breast Cancer Family Registry. *Cancers (Basel)*. Mar 18 2021; 13(6). PMID 33803639

30. Rainville I, Hatcher S, Rosenthal E, et al. High risk of breast cancer in women with biallelic pathogenic variants in CHEK2. *Breast Cancer Res Treat.* Apr 2020; 180(2): 503-509. PMID 31993860
31. Lu HM, Li S, Black MH, et al. Association of Breast and Ovarian Cancers With Predisposition Genes Identified by Large-Scale Sequencing. *JAMA Oncol.* Jan 01 2019; 5(1): 51-57. PMID 30128536
32. Kurian AW, Hughes E, Handorf EA, et al. Breast and Ovarian Cancer Penetrance Estimates Derived From Germline Multiple-Gene Sequencing Results in Women. *JCO Precis Oncol.* Nov 2017; 1: 1-12. PMID 35172496
33. Fan Z, Ouyang T, Li J, et al. Identification and analysis of CHEK2 germline mutations in Chinese BRCA1/2-negative breast cancer patients. *Breast Cancer Res Treat.* May 2018; 169(1): 59-67. PMID 29356917
34. Hauke J, Horvath J, Gro E, et al. Gene panel testing of 5589 BRCA1/2-negative index patients with breast cancer in a routine diagnostic setting: results of the German Consortium for Hereditary Breast and Ovarian Cancer. *Cancer Med.* Apr 2018; 7(4): 1349-1358. PMID 29522266
35. Decker B, Allen J, Luccarini C, et al. Rare, protein-truncating variants in ATM, CHEK2 and PALB2, but not XRCC2, are associated with increased breast cancer risks. *J Med Genet.* Nov 2017; 54(11): 732-741. PMID 28779002
36. Couch FJ, Shimelis H, Hu C, et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. *JAMA Oncol.* Sep 01 2017; 3(9): 1190-1196. PMID 28418444
37. Nslund-Koch C, Nordestgaard BG, Bojesen SE. Increased Risk for Other Cancers in Addition to Breast Cancer for CHEK2*1100delC Heterozygotes Estimated From the Copenhagen General Population Study. *J Clin Oncol.* Apr 10 2016; 34(11): 1208-16. PMID 26884562
38. Huzarski T, Cybulski C, Wokolorczyk D, et al. Survival from breast cancer in patients with CHEK2 mutations. *Breast Cancer Res Treat.* Apr 2014; 144(2): 397-403. PMID 24557336
39. Kriege M, Hollestelle A, Jager A, et al. Survival and contralateral breast cancer in CHEK2 1100delC breast cancer patients: impact of adjuvant chemotherapy. *Br J Cancer.* Aug 26 2014; 111(5): 1004-13. PMID 24918820
40. Weischer M, Nordestgaard BG, Pharoah P, et al. CHEK2*1100delC heterozygosity in women with breast cancer associated with early death, breast cancer-specific death, and increased risk of a second breast cancer. *J Clin Oncol.* Dec 10 2012; 30(35): 4308-16. PMID 23109706
41. Weidner AE, Liggin ME, Zuniga BI, et al. Breast cancer screening implications of risk modeling among female relatives of ATM and CHEK2 carriers. *Cancer.* Apr 15 2020; 126(8): 1651-1655. PMID 31967672
42. Lee A, Mavaddat N, Wilcox AN, et al. BOADICEA: a comprehensive breast cancer risk prediction model incorporating genetic and nongenetic risk factors. *Genet Med.* Aug 2019; 21(8): 1708-1718. PMID 30643217
43. Hall ET, Parikh D, Caswell-Jin JL, et al. Pathogenic variants in less familiar cancer susceptibility genes: what happens after genetic testing? *JCO Precision Oncology.* 2018; 2: 1-10. DOI: 10.1200/PO.18.00167
44. Cragun D, Weidner A, Tezak A, et al. Cancer risk management among female BRCA1/2, PALB2, CHEK2, and ATM carriers. *Breast Cancer Res Treat.* Jul 2020; 182(2): 421-428. PMID 32445176
45. Moslemi M, Vafaei M, Khani P, et al. The prevalence of ataxia telangiectasia mutated (ATM) variants in patients with breast cancer patients: a systematic review and meta-analysis. *Cancer Cell Int.* Sep 08 2021; 21(1): 474. PMID 34493284
46. Marabelli M, Cheng SC, Parmigiani G. Penetrance of ATM Gene Mutations in Breast Cancer: A Meta-Analysis of Different Measures of Risk. *Genet Epidemiol.* Jul 2016; 40(5): 425-31. PMID 27112364
47. Suszynska M, Kozlowski P. Summary of BARD1 Mutations and Precise Estimation of Breast and Ovarian Cancer Risks Associated with the Mutations. *Genes (Basel).* Jul 15 2020; 11(7). PMID 32679805
48. American College of Radiology (ACR). ACR Appropriateness Criteria: Breast Cancer Screening. 2017. <https://acsearch.acr.org/docs/70910/Narrative/>. Accessed July 19, 2023.
49. The American Society of Breast Surgeons. Consensus Guidelines on Genetic Testing for Hereditary Breast Cancer. 2019. <https://www.breastsurgeons.org/docs/statements/Consensus-Guideline-on-Genetic-Testing-for-Hereditary-Breast-Cancer.pdf>. Accessed July 19, 2023.
50. National Comprehensive Cancer Network (NCCN). NCCN Clinical Practice Guidelines in Oncology: Genetic/Familial High-Risk Assessment: Breast, Ovarian, and Pancreatic. Version 3.2023. https://www.nccn.org/professionals/physician_gls/pdf/genetics_bop.pdf. Accessed July 19, 2023.
51. National Comprehensive Cancer Network (NCCN). NCCN Clinical Practice Guidelines in Oncology: Breast Cancer Screening and Diagnosis. Version 1.2023. https://www.nccn.org/professionals/physician_gls/pdf/breast-screening.pdf. Accessed July 18, 2023.

POLICY HISTORY - THIS POLICY WAS APPROVED BY THE FEP® PHARMACY AND MEDICAL POLICY COMMITTEE ACCORDING TO THE HISTORY BELOW:

Date	Action	Description
December 2023	New policy-FEP	Policy updated with literature review through July 17, 2023; no references added. Removed outdated clinical input. Policy statement unchanged. New FEP Policy.

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