

Electronic Medical Record Alerts to Improve High-risk Breast Cancer Patients' Rate of Referral to Genetic Counseling

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INTRODUCTION

- 10-15% of breast cancer cases are hereditary in nature with BRCA1 and BRCA2 mutations accounting for the majority and increasing risk for various malignancies
- National Comprehensive Cancer Network (NCCN) has set forth guidelines to identify patients meeting criteria for genetic risk evaluation so that further testing and risk reduction strategies may be offered
- The Quality Oncology Practice Initiative cites "referral to genetic counseling for high risk breast cancer patients" as a core quality metric.
- Nationwide referral rates are low, including Leo Jenkins Cancer Center (LJCC), with the average referral rate between 2007-2014 of 59%
- Recent research has shown improvements of referral rates for ovarian cancer patients by using electronic medical records (EMR)

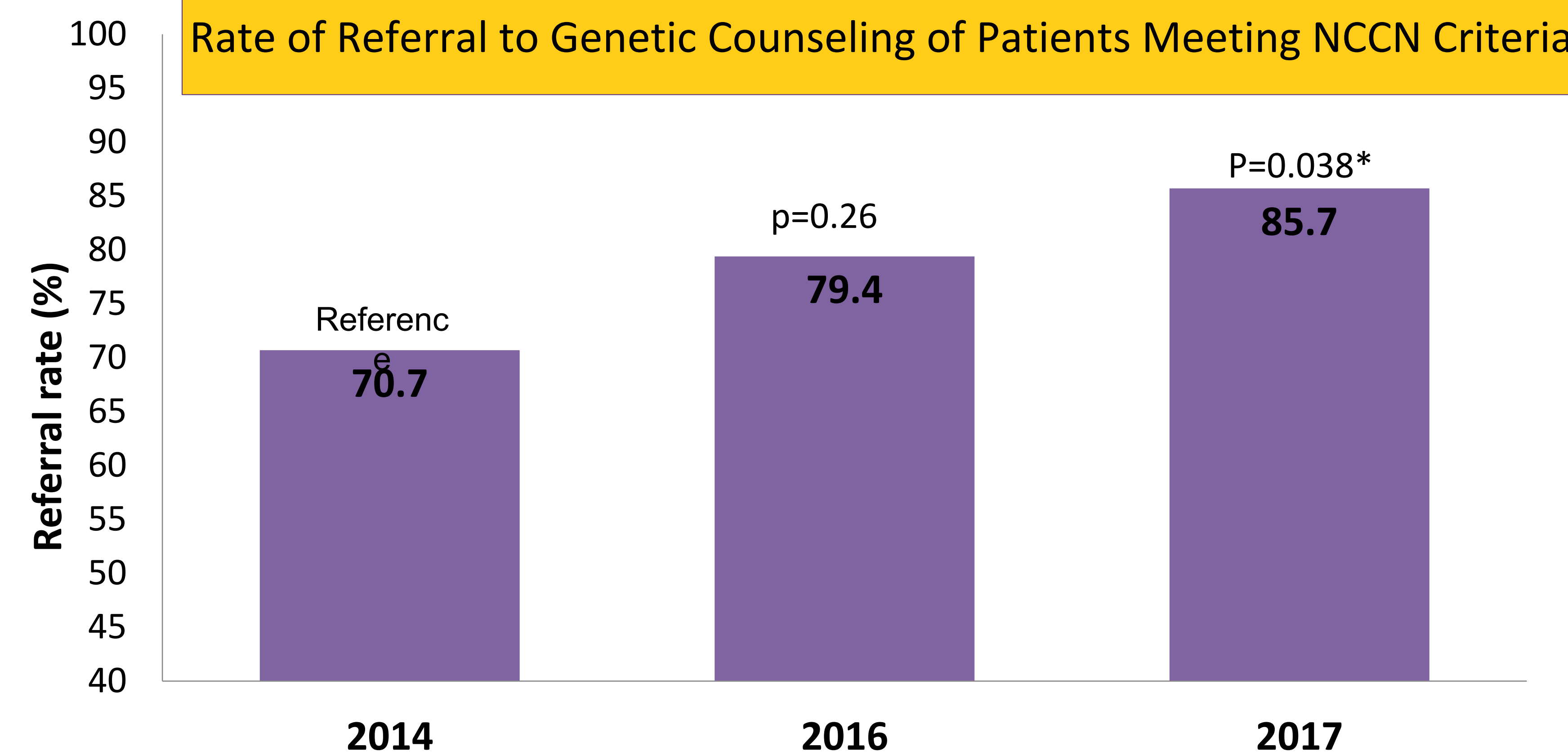
OBJECTIVE

- We sought to assess compliance with NCCN guidelines for genetic risk evaluation and testing among breast cancer patients treated at LJCC
- We hypothesize that implementing an electronic medical record identification and referral system will increase referral rates from 2014 rate of 70% to approximately 80% or higher
- We also wish to increase genetic counseling rates by utilizing nurse navigator follow up to further explore the reasons why patients do not attend counseling

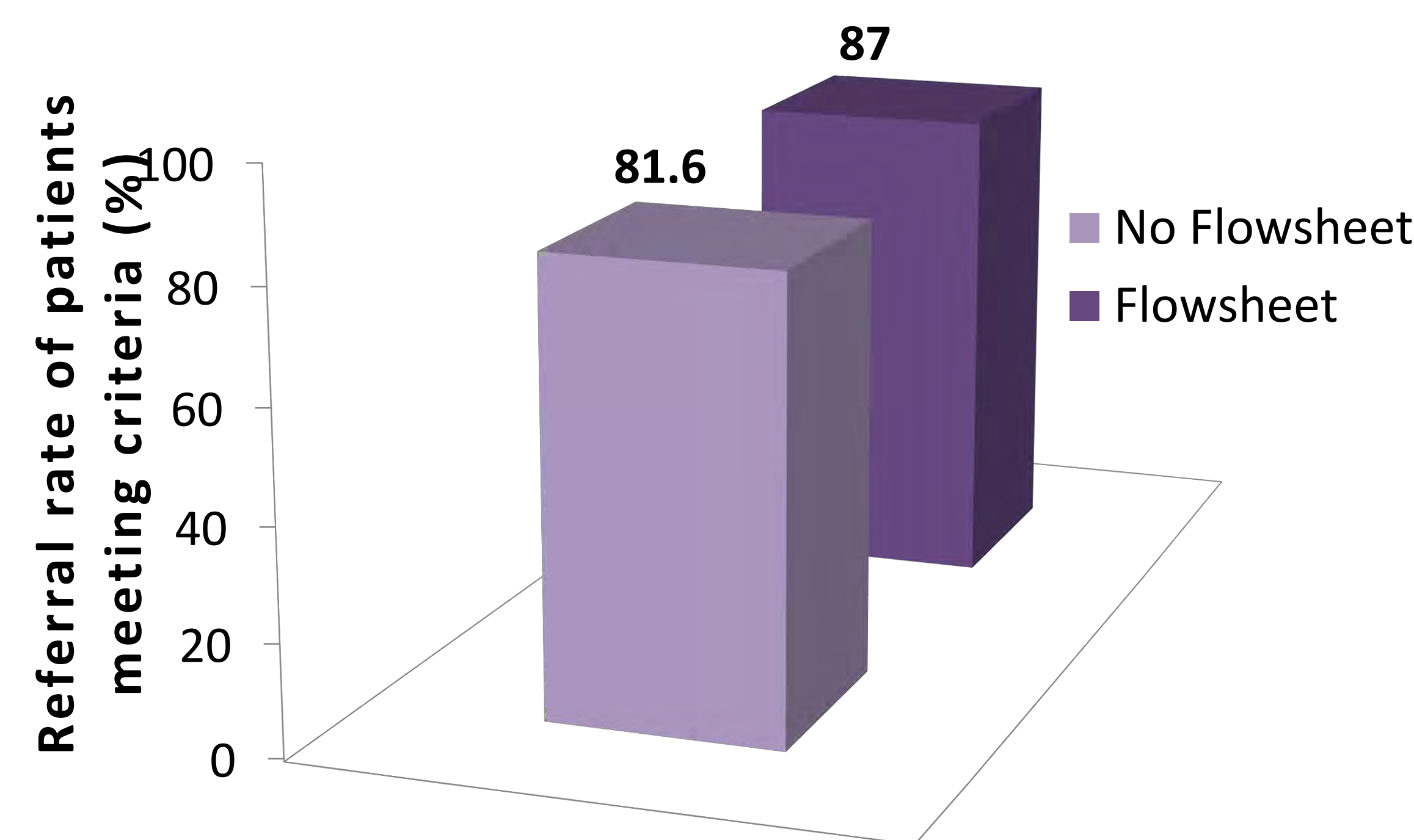
MATERIALS & METHODS

- A document flowsheet was implemented in EPIC at LJCC in 2017 that automatically sends a genetic referral if NCCN criteria are met
- Retrospective electronic chart review of 223 patients diagnosed with breast cancer at LJCC between January 2016 and August 2017, to assess rates of genetics referral, counseling, and testing
- NCCN guidelines were used to determine if patients met criteria for further genetic risk evaluation
- Rates of referral, counseling and testing were determined for those who met criteria using chi-squared and odds ratio
- Pre (2016) and post (2017) EMR template implementation was analyzed
- Reasons for not undergoing genetic counseling were assessed

RESULTS



Improved Genetic Counseling Referral Rate for Patients with Flowsheet Filled



Reason for not undergoing genetic counseling		
Reasons	2016 N(%)	2017 N(%)
Refused	7 (41.1)	4 (22.2)
Financial	1 (5)	3 (16.6)
No Show	1 (5)	5 (27.7)
Undocumented	4 (23.5)	0 (0)

SUMMARY

- A total of 108 genetic document flowsheets were filled by providers
- Of patients diagnosed in 2017, 72.3% of flowsheets were filled
- The flow sheet was significantly more likely to be filled if the patient met criteria (56%) compared to those who did not meet criteria (36%)
- Referral rates showed an increasing trend with time
- In 2017, for patients who met NCCN criteria and had a genetic flowsheet completed, the referral rate was 90.6%
- Genetic counseling and testing rates remained the same with time
- Reasons for not undergoing counseling were better documented in 2017, with none unknown compared to 23% unknown in 2016
- Patients from counties with low median household income were less likely to attend genetic counseling once referred

CONCLUSIONS

- Improvements have been seen in referral rates of high-risk breast cancer patients after utilizing the EMR flowsheets
- This difference was especially large for those who met the less obvious criteria, family history of pancreatic and/or prostate cancer (22% in 2010-2014 vs 91% in 2016-2017)
- About three quarters of flowsheets were filled in 2017, so efforts must be continued in order to reach 100%
- Ten patients who met criteria and had a flowsheet completed were not referred for genetic risk assessment; this was determined to represent a technical issue of the automatic referral not being triggered, which has since been resolved
- Despite increased rates of referral, rates of attending counseling did not significantly increase, so patient education and nurse navigator follow up are increasingly important

FUTURE DIRECTIONS

- We propose using similar flowsheets across the Vidant Health Network oncology practices for all genetic cancers