

Back to the future: pathology

MOLECULAR DRIVEN ONCOLOGY: FUTURE DIRECTIONS

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Disclosure of Interest

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Nature, 2015 source: Pharmaceutical Research and Manufacturers of America FTLO Science Washington DC: Foundation For Biomedical Research Wong, C. Biostatistics, 2018. Mestre-Ferrandiz, J.2012 Paul S.M., et al, Nature reviews Drug discovery 2010







Genomics in healthcare: clinical utility





Clinical utility: target therapies





J. Mateu et al, Nature 2022

Clinical utility: the case of ovarian cancer





Jdi Silvestro et al, JCO 2023

Clinical utility: the case of endometrial cancer





Bokhman, Gynecol Oncol 1983 TGCA, Nature 2013 Stelloo, Clin Can Res 2016

Clinical utility: the case of endometrial cancer

Table 2. EC risk groups	
Risk group	Description ^a
Low risk	Stage IA (G1-G2) with endometrioid type (dMMR ^b and NSMP) and no or focal LVSI Stage I/II POLEmut cancer; for stage III POLEmut cancers ^c
Intermediate risk	Stage IA G3 with endometrioid type (dMMR and NSMP) and no or focal LVSI Stage IA non-endometrioid type (serous, clear-cell, undifferentiated carcinoma, carcinosarcoma, mixed) and/o <mark>r p53-abn</mark> cancers without myometrial invasion and no or focal LVSI Stage IB (G1-G2) with endometrioid type (dMMR and NSMP) and no or focal LVSI Stage II G1 endometrioid type (dMMR and NSMP) and no or focal LVSI
High-intermediate risk	Stage I endometrioid type (dMMR and NSMP) any grade and any depth of invasion with substantial LVSI Stage IB G3 with endometrioid type (dMMR and NSMP) regardless of LVSI Stage II G1 endometrioid type (dMMR and NSMP) with substantial LVSI Stage II G2-G3 endometrioid type (dMMR and NSMP)
High risk	All stages and all histologies with <mark>p53-abn a</mark> nd myometrial invasion All stages with serous or undifferentiated carcinoma including carcinosarcoma with myometrial invasion All stage III and IVA with no residual tumour, regardless of histology and regardless of molecular subtype ^b



Oaknin A. et al Ann Oncol 2022





PREVENTION IS BETTER THAN EARLY DETECTION

Genomics profiling as an opportunity for cascade prevention



- **13.3%** PGVs found using **CGP**
- **48.4%** of them would not have been detected by phenotype or family history-based testing criteria using the 2018 NCCN, NSGC, or ACMG guidelines
- **17.6%** with PGVs had family members undergoing no-cost cascade FVT



- **2.22%** PGVs in in moderate risk
 breast and ovarian cancer susceptibility
 genes found using CGP
- **45.3%** of them would not have eligible for germline testing
- **51.5%** of the ones undergoing germline testing (40% of the identified) had germline confirmation



- O 11.2% PVGs found using CGP
- 9.2% PVGs found using standard approaches



N.J. Samadder et al., JAMA oncol 2021 Llorin H et al, Cancer Genet. 2022 N. Normanno et al, EJC 2023

TAILORING DOSE AND TREATMENTS

Pharmacogenomics and genomic-adjusted radiation dose



MODERN RADIATION ONCOLOCY



M. Pirmohamed,et al.,. Nature Reviews Genetics 2023 Scott JG et al., Lancet Oncol. 2017

DECISION MAKING

The role of liquid biopsy in minimal residual disease assessment









TORWARDS PERSONALIZED MEDICINE

Integrated Multi-omics longitudinal profiling





TORWARDS PERSONALIZED MEDICINE

From data to information







CHALLENGES

Complex data analysis and integration

Molecular diagnostic market, Regional Analysis 2021-2031 from Fact.MR Kovanda et al., Human Genomics 2021

MOLECULAR DRIVEN ONCOLOGY

Inevitable evolution

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