

NGS tests and algorithms in (hemato)-oncology

[Symposium NGS 2016 | sciensano.be](https://www.sciensano.be)
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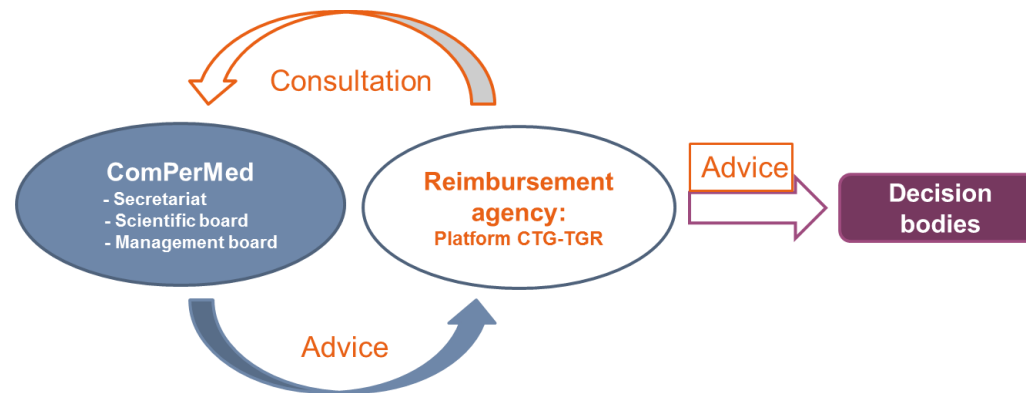
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Introduction

- Targeted NGS is gradually introduced in molecular diagnostics in daily practice as it enables more precise decision-making:
 - diagnosis
 - prognosis
 - therapeutic choices for targeted drugs or conventional treatments
 - A permanent monitoring and assessment of the state-of-the-art in this domain is essential
- Commission of Personalized Medicine (ComPerMed)

Commission Personalized Medicine (ComPerMed)

- **Aim:** provide evidence-based advices to the platform CTG-TGR of the RIZIV-INAMI
 - technical aspects of NGS testing
 - clinical use of NGS for detecting somatic mutations in cancer patients
 - assessment of novel 'omics' technologies



- Belgian professionals and experts in the field of (hemato)-oncology

Technical aspects of NGS

NGS guidelines:

- to **facilitate the implementation** of NGS in the laboratories
- to help labs to **generate accurate NGS data**
e.g. Identical sample analysis should lead to an identical list of variants even if processed by a different operator on a different day.
- to **facilitate the evaluation by the auditors** from the accreditation bodies (Belac)



The Belgian next generation sequencing guidelines for haematological and solid tumours

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Clinical use of NGS

Selection of genes with regard to their **test utility** for a specific tumor

→ **diagnosis**

→ **therapeutic** response (to predict **sensitivity** or **resistance**)

→ **prognosis** (patient outcome)



biomarker/test levels

Level 1
Level 2
Level 3

- Standard of care biomarker for diagnosis and/or prognosis *
- Biomarker predictive of a response or a resistance to a reimbursed drug in Belgium for this indication

A

- Recommended standard of care biomarker for diagnosis and/or prognosis +
- Biomarker predictive of response or resistance to an EMA-approved drug for this indication

B

- Biomarker predictive of response or resistance to a reimbursed drug in Belgium for another indication (clinical trial available in Belgium or EU)

- Compelling clinical evidence supporting the biomarker for diagnosis and/or prognosis
- Biomarker predictive of a response or a resistance to
 - a non EMA-approved drug in this indication
 - a reimbursed drug in Belgium for another indication (clinical trial not available in Belgium or EU)
 - an EMA-approved drug for another indication
- Compassionate use of drug

* Standard of care: Included in guidelines (WHO) AND consensus from experts ComPerMed

+ Recommended standard of care: Clinical evidence AND consensus from experts ComPerMed

Level 1

- Standard of care biomarker for diagnosis and/or prognosis *
- Biomarker predictive of a response or a resistance to a reimbursed drug in Belgium for this indication

Standard of care

Level 2

A

- Recommended standard of care biomarker for diagnosis and/or prognosis +
- Biomarker predictive of response or resistance to an EMA-approved drug for this indication

Level 3

B

- Biomarker predictive of response or resistance to a reimbursed drug in Belgium for another indication (clinical trial available in Belgium or EU)

**Clinical trial
compassionate use
research**

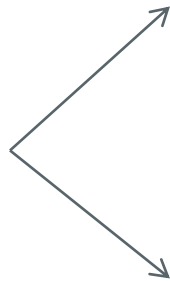
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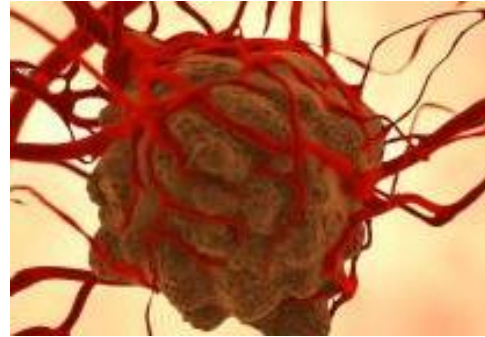
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Clinical use of NGS

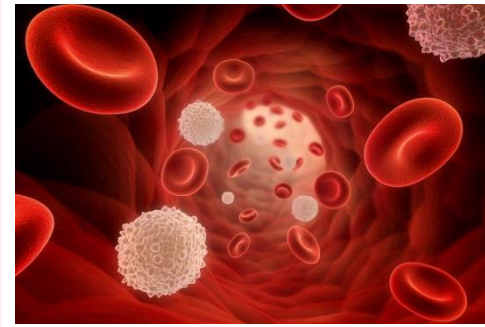
Test utility and level



Solid tumors

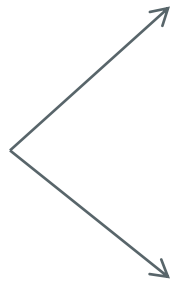


Hematological tumors



Clinical use of NGS

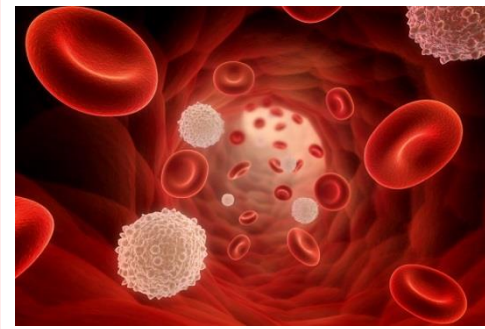
Test utility and level



Solid tumors



Hematological tumors

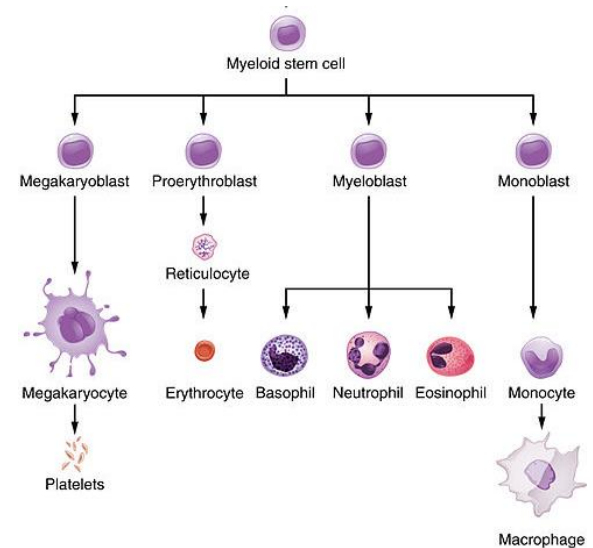


MYELOID CANCERS

Clinical use of NGS

List of myeloid cancers for which an NGS test is useful:

- Acute myeloid leukemia (AML)
- Myeloproliferative neoplasm (MPN)
 - Chronic neutrophilic leukemia (CNL)
 - Primary Myelofibrosis (PMF)
- Myelodysplastic syndrome (MDS)
- MDS/MPN
 - Chronic myelomonocytic leukemia (CMML)
 - Atypical CML (aCML)
 - MDS/MPN-RS-T



Clinical use of NGS



List of genes which should be analyzed at minimum by NGS: level 1 & 2A

Acute myeloid leukemia			
Genes	Diagnosis	Prognosis	Therapy
ASXL1	x (2A)	x (1)	
CEBPA*	x (1)	x (1)	
DNMT3A*	x (2A)	x (1)	
FLT3	x (2A)	x (1)	x (2B)
IDH1	x (2A)	x (1)	x (3)
IDH2	x (2A)	x (1)	x (3)
KIT	x (2A)	x (1)	x (2B)
NPM1	x (1)	x (1)	
RUNX1*	x (1)	x (1)	
TET2*	x (2A)	x (1)	
TP53*	x (2A)	x (1)	x (3)
WT1	x (2A)	x (1)	

Clinical use of NGS



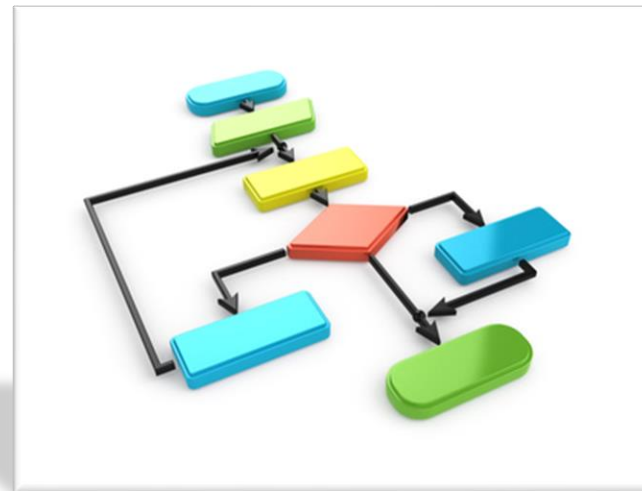
List of genes which should be analyzed at minimum by NGS: level 1 & 2A

Genes	AML	MDS	MPN PMF	MPN CNL	MDS/MPN aCML	MDS/MPN CMML	MDS/MPN RS-T
ASXL1	x	x	x	x	x	x	
CALR			x		x		x
CEBPA	x						
CSF3R				x	x		
DNMT3A	x	x					
EZH2		x	x				
FLT3	x						
IDH1	x		x				
IDH2	x		x				
JAK2			x		x		x
KIT	x						
MPL			x		x		x
NPM1	x						
RUNX1	x	x					
SETBP1				x	x	x	
SF3B1		x	x				x
SRSF2		x	x	x	x	x	
TET2	x	x	x	x		x	
TP53	x	x					
U2AF1		x					
WT1	x						

Clinical use of NGS

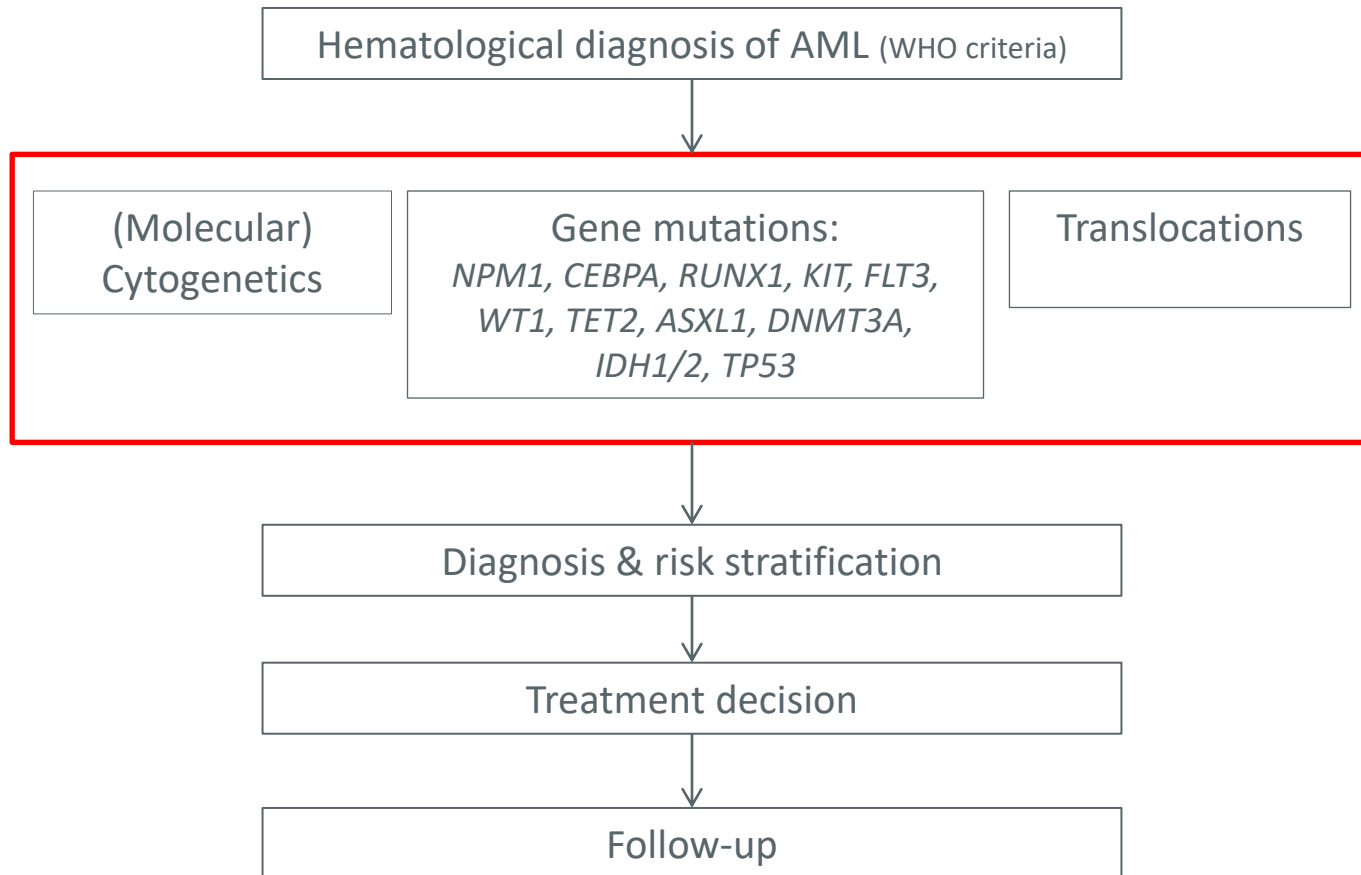
Algorithms

- Display the NGS test in the context with all the other molecular tests (FISH, IHC, PCR,...)
- Define the indications for NGS testing



Clinical use of NGS

Example: algorithm of acute myeloid leukemia



Clinical use of NGS

Integrate NGS testing and algorithm in a website



Clinical use of NGS



Integrate NGS testing and algorithm in a website

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Algorithms

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Incidence
List of experts
Algorithms

```
graph TD; A[All CRC] --> B[TEST: mismatch repair (MMR) proteins (MLH1, MSH2, MSH6, PMS2) (IHC)]; B --> C[TEST: PCR MSI]; C --> D[IF Loss of MLH1 AND PMS2 (MSI-H)]; C --> E[IF Loss of MSH2 MSH6 OR PMS2 (without loss of MLH1) (MSI-H)]; C --> F[IF No loss of nuclear expression for MMR proteins (MSS AND/OR MSI-L)]; D --> G[TEST: mut BRAF]; G --> H[If TEST +]; G --> I[If TEST -]; F --> J[Family history suspect for Lynch syndrome / or other familial cancer syndromes?]; J --> K[IF Yes]; J --> L[IF No];
```

The flowchart details the following steps:

- Start:** All CRC
- Test 1:** TEST: mismatch repair (MMR) proteins (MLH1, MSH2, MSH6, PMS2) (IHC). Levels of evidence: 1. Test description.
- Test 2:** TEST: PCR MSI
- Branch 1:** IF Loss of MLH1 AND PMS2 (MSI-H) → TEST: mut BRAF → If TEST + / If TEST -
- Branch 2:** IF Loss of MSH2 MSH6 OR PMS2 (without loss of MLH1) (MSI-H)
- Branch 3:** IF No loss of nuclear expression for MMR proteins (MSS AND/OR MSI-L) → Family history suspect for Lynch syndrome / or other familial cancer syndromes? → IF Yes / IF No

Clinical use of NGS



Integrate NGS testing and algorithm in a website

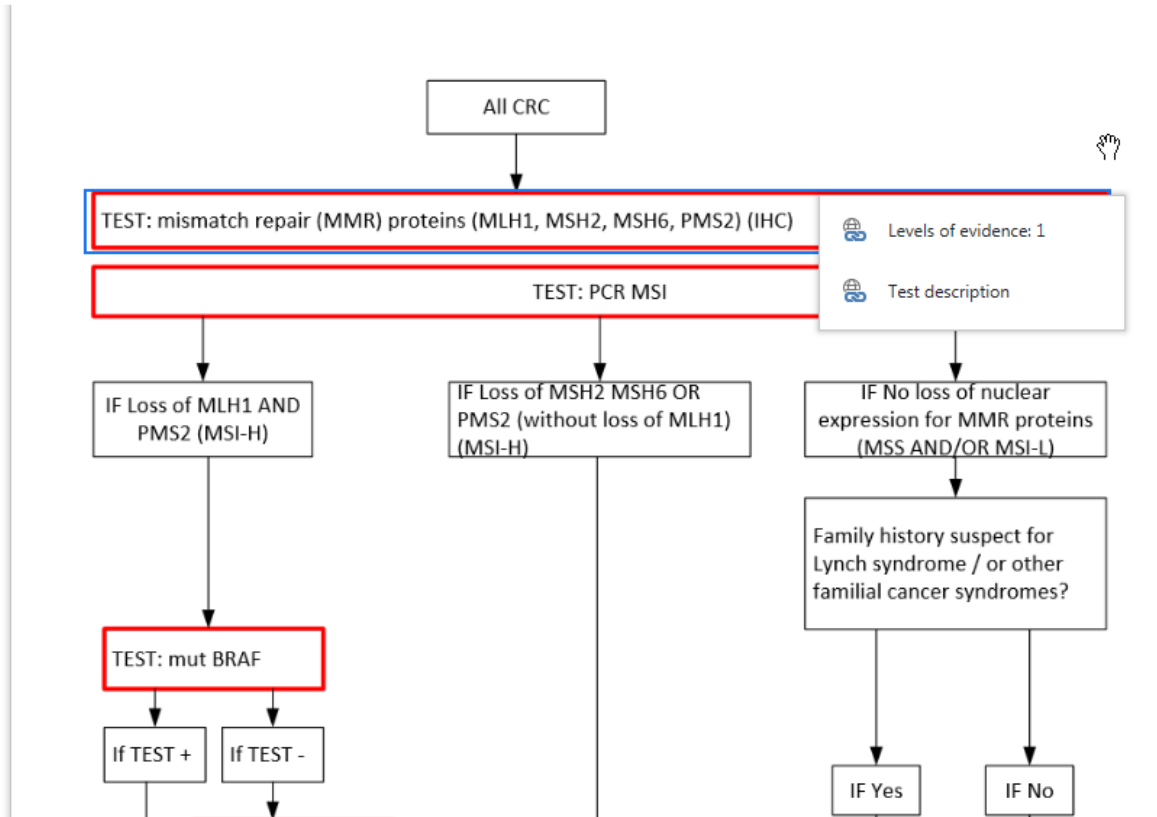
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Algorithms

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- Incidence
- List of experts
- Algorithms



Clinical use of NGS



Integrate NGS testing and algorithm in a website

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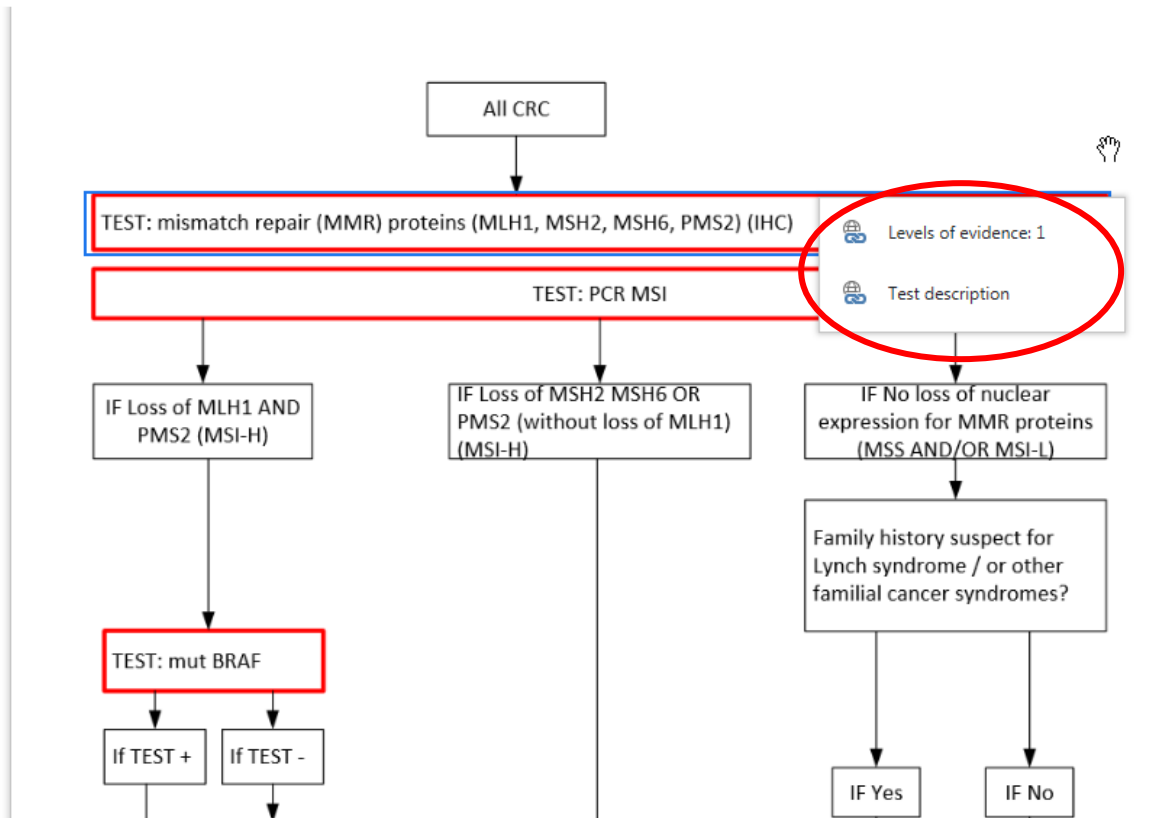
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Incidence

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Clinical use of NGS

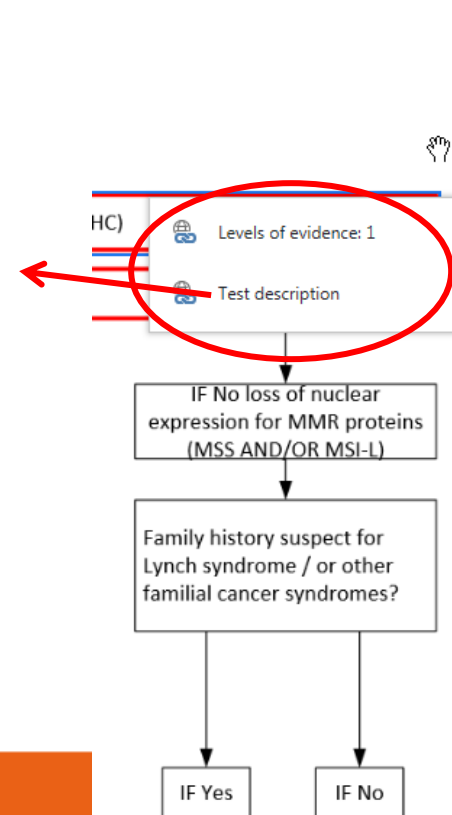
Integrate NGS testing and algorithm in a website

MMR proteins test by IHC and/or PCR MSI in CRC

Microsatellite instability (MSI) is a hypermutable phenotype caused by the loss of DNA mismatch repair activity. Microsatellites are non-coding DNA sequences constituted by repeated units of 2-9 base pairs. They are randomly dispersed throughout the human genome.

Microsatellite instability (MSI) is defined as a change (deletion or insertion of repeating units) in the microsatellite sequences in a tumor compared to the normal tissue. Microsatellite instability (MSI) is caused by a defect in the DNA mismatch repair (MMR) genes which normally occurs to correct errors during DNA replication.

The PCR detection by the amplification of 5 mono/di-nucleotide microsatellite markers is the standard test. MSI tests can be divided into three groups: MSI-H, when $\geq 30\%$ of markers exhibit



Future work

- Website: January 2018
- Regularly update clinical use of NGS: genes and algorithms
- Clinical use of NGS in other hematological cancers (lymphoid)
- Clinical use of NGS in pediatric cancers
- Harmonize variant annotation and clinical interpretation



THANK YOU

