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Molecular pathology in routine diagnostics of solid tumors

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Molecular pathology in routine diagnostics of solid tumors

A. Types of molecular markers

B. Different molecular tools

C. Clinical utility of tumor markers in oncology: examples of molecular testing in routine daily practice

Types of molecular markers

- **DIAGNOSTIC** markers
- **PROGNOSTIC** markers
- **PREDICTIVE** markers

Diagnostic markers

- Large category of molecular tests aid in the diagnosis or subclassification of a particular disease state
- Diagnostic subclassification may result in different management of the disease
especially in soft tissue pathology and hematopathology
 - MDM2 and CDK4 amplification in well-differentiated and dedifferentiated liposarcomas
 - MYC amplification in postirradiation angiosarcoma
 - rearrangement of CHOP, EWS and SY7 genes in myxoid liposarcoma, Ewing sarcoma and synovial sarcoma respectively
 - KIT and PDGFRα mutation in Gastrointestinal stromal tumors (GIST)
 - rearrangement of BCL2 gene in follicular lymphoma
 - rearrangement of MYC gene in Burkitt lymphoma
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Prognostic markers

- Prognostic markers have an association with some clinical outcomes (overall survival, recurrence-free survival), independent of the treatment rendered
e.g. presence of *p53* mutations, which identify subsets of patients who will have a more aggressive disease course for certain cancers, regardless of current treatment options

Predictive markers

- Predictive markers predict the activity of a specific class or type of therapy, and are used to help make more specific treatment decisions
- They are used as indicators of the likely benefit of a specific treatment to a specific patient ("personalized medicine")
e.g. *HER2* status in breast and gastric cancer: *HER2*-positivity (amplification) is predictive of potential trastuzumab response
e.g. *BRAF V600E* mutation is predictive of sensitivity to vemurafinib

Molecular diagnostic tools

- **conventional cytogenetic/chromosome analysis**
short-term, primary cultures followed by karyotyping

Conventional karyotyping

- **Advantages:**
 - global genetic information in single assay
 - variants uncovered (undetectable by FISH and RT-PCR)
 - diagnostically useful: sensitive, specific
 - provides direction for further molecular studies
- **Limitations:**
 - requires fresh tissue
 - mostly cell culture 1-10 days
 - complex karyotypes, suboptimal morphology
 - normal karyotypes (overgrowth normal fibroblasts, infiltrating cells)

Molecular diagnostic tools

- **Molecular cytogenetics**
 - * **fluorescence in situ hybridization (FISH):** detecting of cancer-related translocations, deletions and amplifications
 - * **molecular karyotyping (cytogenomic arrays) (CGH array, SNP analysis):** detecting whole genome copy number changes including losses, gains and amplifications

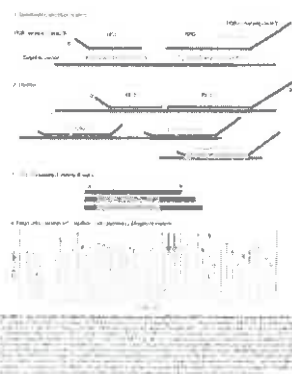
Molecular diagnostic tools

- **Molecular assays**
 - * **Polymerase chain reaction (PCR), reverse transcription-polymerase chain reaction (RT-PCR):** assays for known mutations, translocations
 - * **Multiplex Ligation-dependent Probe Amplification (MLPA):** detecting losses, gains and amplifications
 - * **Expression profiling (cDNA arrays)**
 - * **Next generation whole genome technologies (NGS)**

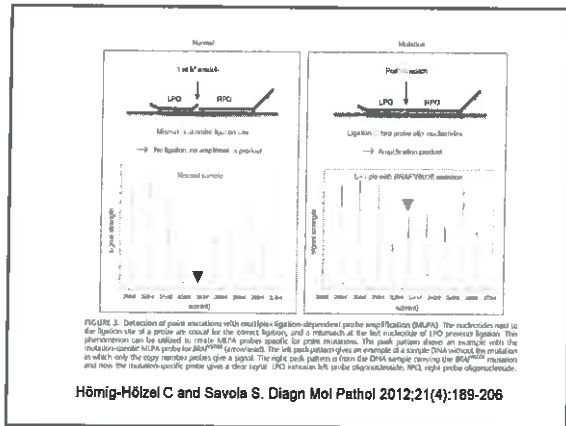
Multiplex Ligation-dependent Probe Amplification

- Special form of **multiplex PCR**
- Detection of aberrant **copy number** of 55 genomic DNA sequences in a single, PCR-based, reaction
- MLPA can also be used for detecting **methylation status** and **known point mutations**
- **Amplification of MLPA probes, not sample DNA!**
- Requires a minimum of 20ng of human DNA (3000 cells)
- MLPA can also be used on partially degraded DNA (e.g. FFPE)

Hörig-Hötzel C and Savola S. *Diagn Mol Pathol* 2012;21(4):189-206
 Creytens D, van Gorp J, Ferdinande L, Speel EJ and Libbrecht L. *Diagn Mol Pathol* 2014
 2014;46(5(1)):37-108
 Creytens D, van Gorp J, Speel EJ and Ferdinande L. *Anticancer Res* 2014



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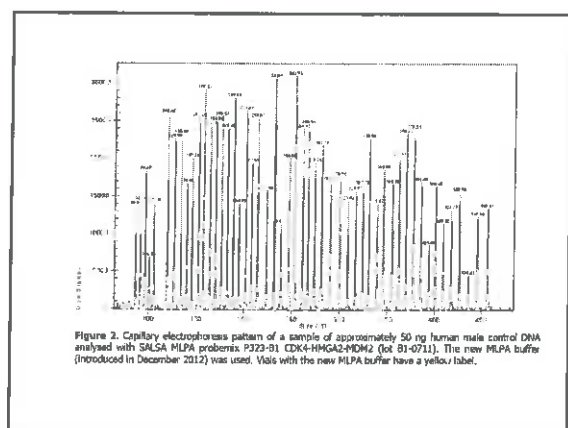
- ### Advantages of MLPA for tumor applications
- MLPA is excellent technique for large scale validation of new biomarkers and for use in routine diagnostics
 - Probe for both methylation and copy number changes and a few probes for specific point mutations can be combined in one test
 - High throughput: up to 96 samples can be analyzed in 24h, with little hands on time; 55 targets can be analyzed simultaneously!

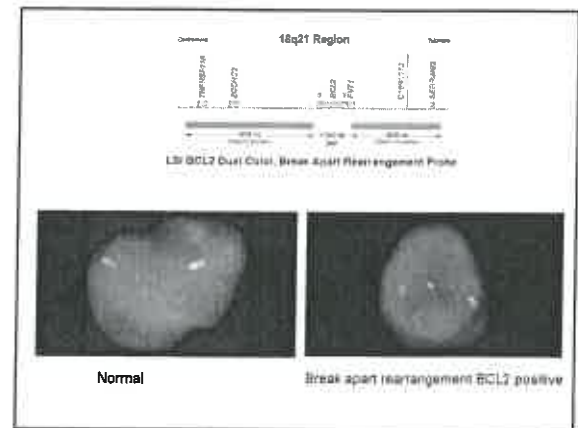
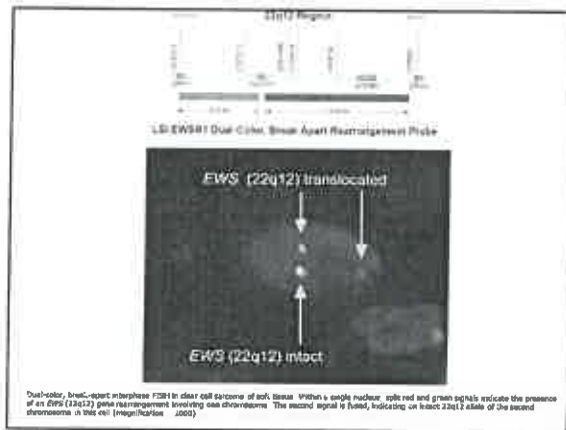
- ### Limitations of MLPA for tumor applications
- MLPA detects in RELATIVE amounts of target sequences: gains/losses may not be detected if percentage of tumor cells is too low or due subclonality (<20-30%)
 - NO MORPHOLOGICAL CORRELATION
 - Selection of reference probes for MLPA mix is critical: they should locate on "silent regions" of the tumor genome
 - MLPA technology is not suitable for the detection of unknown mutations

- ### MLPA
- **Copy number detection:** *MYCN* (neuroblastoma), *HER2* (breast cancer, gastric cancer), *MDM2* (liposarcoma), *EGFR* (glioblastoma)
 - **Point mutation detection with MLPA*:** *IDH1/2* (diffuse glioma), *BRAF* (melanoma, pilocytic astrocytoma), *EGFR* (lung carcinoma)
- *genes with less-defined hot spots or many point mutations within a short distance from each other (TP53, p16/CDKN2A), are less suitable for the design of mutation-specific MLPA probes!

Group	Gene	MLPA probe	Chromosomal location
1	11p15.5	11p15.5	11p15.5
	12p12	12p12	12p12
	13q32	13q32	13q32
	14q32	14q32	14q32
	15q21	15q21	15q21
	16p11	16p11	16p11
	17q21	17q21	17q21
	18q21	18q21	18q21
	19p13	19p13	19p13
	20p12	20p12	20p12
2	21q22	21q22	21q22
	22q11	22q11	22q11
	22q13	22q13	22q13
	22q14	22q14	22q14
	22q15	22q15	22q15
	22q16	22q16	22q16
	22q17	22q17	22q17
	22q18	22q18	22q18
	22q19	22q19	22q19
	22q20	22q20	22q20

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FISH: advantages

- Fresh, frozen, paraffin-embedde material
- Localize alteration (amplification, translocation) in specific cells an tissue types)
- Useful if tumor is heterogeneous
- Diagnostically useful: fine needle biopsy, sensitive, specific
- Can provide results if karyotyping or RT-PCR is inconclusive
- Rapid turn-around time
- Validation and implemenation easy
- Normal tissue parts can serve as FISH control

FISH: limitations

- Relatively gross approach (no information on fusion genes and variants)
- Number of commercially available probes is limited (Abbott Molecular, Kreatech)
- Requires fluorescence microscope
- Interpretation may be challenging, expertise required
- Period of storage

