

The evolving role of molecular techniques in surgical pathology:

focus on soft tissue and bone tumour diagnostics

Fleur Cordier

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Ghent University Hospital

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Curriculum vitae

Qualifications

Medical Doctor, 2018, KU Leuven, Belgium
Pathologist, 2023, Ghent University, Belgium
(Co-) author of 28 publications in peer reviewed
international journals

Main publications PhD

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tumours. J Clin Pathol. 2023

Promoter

Prof. Dr. David Creyten
Department of Diagnostic Sciences and Department of
Pathology
Ghent University and Ghent University Hospital, Belgium

Co-promoter

Prof. Dr. Jo Van Dorpe
Department of Diagnostic Sciences and Department of
Pathology
Ghent University and Ghent University Hospital, Belgium

Doctoral Examination Committee

Prof. Dr. Patrick Calders (Chair)
Department of Rehabilitation Sciences and Physiotherapy
Ghent University, Belgium

Prof. Dr. Lore Lapeire

Department of Internal Medicine and paediatrics and
Department of Medical Oncology
Ghent University and Ghent University Hospital, Belgium

Prof. Dr. Olivier De Wever

Laboratory of Experimental Cancer Research
Department of Human Structure and Repair
Ghent University, Belgium

Prof. Dr. Raf Sciôt

Department of Pathology
University Hospitals Leuven, Belgium

Dr. Vasiliki Siozopoulou

Department of Pathology
Saint-Luc University Hospital, Brussels, Belgium

Prof. Dr. Gwen Sys

Department of Human Structure and Repair and
Department of Orthopaedics and Traumatology
Ghent University and Ghent University Hospital, Belgium

Prof. Dr. Isabelle Vanden Bempt

Department of Human Genetics
University Hospitals Leuven, Belgium

Funding: Pathology Department, Ghent University Hospital

Summary

The implementation of (new) molecular techniques in the daily practice has transformed the role of pathologists, transitioning from a morphologic focus to a comprehensive morphomolecular perspective, encapsulated in the concept of 'integrated diagnostics'. Recent years have witnessed a rapid integration of advanced molecular techniques like next-generation sequencing (NGS) into routine pathology practices. This has significantly enhanced our comprehension of disease pathogenesis and led to the identification of novel tumour entities or subtypes. However, with this wealth of information comes the challenge of navigating through it.

Molecular findings challenge traditional taxonomies, urging us to reconsider the boundaries of diseases and the concept of distinct entities. Entering the realm of the 'splitters' and the 'lumpers', the 'splitters' advocate for finer distinctions based on molecular signatures, while the 'lumpers' argue for overarching categories that encompass shared characteristics. This debate surfaces in this thesis in which 2 projects were conducted.

1. '*USP6-associated neoplasms*' (*UANs*): Despite a common genetic pathway with *USP6* rearrangements leading to *USP6* overexpression, morphological and clinical differences persist among entities like aneurysmal bone cyst (ABC) and nodular fasciitis (NF). In an exhaustive study of 175 cases using RNA NGS and fluorescence in situ hybridization (FISH) a significant entity- and location-specific *USP6* fusion partner diversity

was revealed. We suggested that observed variations between members of the *UAN* are attributable to a distinct cell of origin. This suggests that these lesions can be classified as distinct entities within the *UAN* group.

2. '*RB1*-deleted soft tissue tumours': We identified the loss of *RB1*, using FISH, in fibroepithelial stromal polyps and in two cases of *PRRX*-rearranged mesenchymal tumours, suggesting an expansion of the group of '*RB1*-deleted soft tissue tumours'. However, controversies surrounding the FISH cut-off persist and warrants further exploration.

Furthermore, we elucidate that the use of molecular techniques not only uncovers new gene rearrangements and tumour entities but also proves invaluable in confirming diagnoses in uncommon locations. In some instances, making a diagnosis without these molecular techniques would be impossible. The discovery of new genetic alterations also leads to the development of surrogate immunohistochemical markers. However, the correlations between these markers and the underlying gene rearrangements are not always straightforward.

This thesis further discusses various aspects including epidemiology, clinical practice, and artificial intelligence. Yet, the fundamental question remains:

Should we split or lump?

CONTACT

Department of Diagnostic Sciences Ghent
Department of Pathology
Ghent University Hospital
Corneel Heymanslaan 10 9000 Gent

fleur.cordier@ugent.be

An electronic version of this doctoral thesis
can be requested by email

T +32 9 332 6782

www.ugent.be