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| Teacher(s)          | Vikkula Miikka ;  |
| Language :          | French<br>> English-friendly  |
| Place of the course | Bruxelles Woluwe  |
| Prerequisites       | <i>The prerequisite(s) for this Teaching Unit (Unité d'enseignement – UE) for the programmes/courses that offer this Teaching Unit are specified at the end of this sheet.</i>  |
| Learning outcomes   |   |
| Evaluation methods  | <p>The student should show his-her acquired knowledge in following situations :<br/>during the exam (multiple choice questions).</p> <p>This assessment is based on a series of multiple-choice questions covering all the material given during the course - 5 proposals / 2 only correct - no negative point. If the student has chosen 0 or only 1 of the correct answers, or more than 2 answers, no point is awarded. Scores are rounded up to the next highest grade if &gt; or equal to 0.5 (e.g. 12,5 becomes 13), except for grades between 9.5 and &lt;10 which are rounded to 9.</p> <p>The book Pasternak contains questions and the book Thompson &amp; Thompson questions and answers for self-learning after each chapter.</p>   |
| Teaching methods    | Teaching is based on lectures (total 20 hours). It relies on the development of theoretical concepts, but also on the description of concrete examples of genetic diseases and genetic analyses.  |
| Content             | <p><b>1. DNA - carrier of genetic information</b></p> <p><b>2. Cytogenetics: low-resolution genome analysis</b></p> <p>2.1. Normal karyotype</p> <p>2.2. Abnormal karyotype</p> <p><b>3. Types of polymorphisms</b></p> <p>3.1. Vocabulary: polymorphism, gene, allele, homozygous, heterozygous</p> <p>3.2. Genome in a single (&lt;10) copy (haploid genome)</p> <p>3.3. Genome moderately repetitive</p> <p>3.4. Genome strongly repetitive</p> <p>3.5. Other polymorphisms</p> <p>3.6. Counting of heterozygosity</p> <p><b>4. Methods to detect genetic markers (polymorphisms)</b></p> <p>4.1. Southern blot</p> <p>4.2. PCR amplification</p> <p>4.3. Molecular karyotyping (DNA arrays)</p> <p>4.4. Sequencing</p> <p>4.5. Next Generation Sequencing (NGS)</p> <p><b>5. Transmission of hereditary characters (Laws of Mendel)</b></p> <p>5.1. How to draw a genealogic tree in genetics - symbols</p> <p>5.2. Types of heredity</p> <p>5.3. Independent and non-independent segregation</p> <p><b>6. Use of polymorphisms</b></p> <p>6.1. Genetic maps</p> <p>6.2. Identification of an individual</p> <p>6.3. Linkage analysis</p> <p>6.4. Autozygosity analysis</p> <p>6.5. Association studies</p> <p>6.6. Loss-of-heterozygosity analyses</p> <p><b>7. Other parameters of heredity</b></p> <p>7.1. <i>de novo</i> mutations</p> <p>7.2. Factors to modify the phenotype</p> <p>7.3. Law of Hardy-Weinberg</p> <p>7.4. eQTL</p> |

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|                             | 7.5. Personalised medicine  |
| Bibliography                | <p><a href="#">Dian-Donnai-Genetique-medicale</a>; De la biologie à la pratique clinique; de Boeck, 2009.<br/>                 New Clinical Genetics 4 (Reed &amp; Donnai)<br/>                 Robert L. Nussbaum, Roderick R. McInnes, Huntington F. Willard. Genetics in Medicine<br/>                 Editeur: Thompson &amp; Thompson, 8e édition, Elsevier, (2016)<br/>                 - Syllabus (Notes de cours vérifié)(iCampus)<br/>                 - Dias du cours (iCampus)<br/>                 et<br/>                 Jack Pasternak : Génétique moléculaire humaine : une introduction aux mécanismes des maladies héréditaires<br/>                 Editeur: De Boek (2003) <b>Chapitres: 1-3(p.3-80), 5.1, 5.4, 5.6, 5.7, 6 (p.161-183)</b></p> |
| Faculty or entity in charge | MED   |

| <b>Programmes containing this learning unit (UE)</b> |                         |         |  |   |
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| Program title  | Acronym                 | Credits | Prerequisite   | Learning outcomes   |
| Bachelor in Medecine                                 | <a href="#">MD1BA</a>   | 2       | <a href="#">WMDS1109</a>   |  |
| Bachelor in Biomedicine                              | <a href="#">SBIM1BA</a> | 2       | <a href="#">WFARM1221S</a> AND <a href="#">WSBIM1226</a><br>AND <a href="#">WMDS1230</a> AND <a href="#">WFARM1247</a> |  |