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| General Information | MASTER DEGREE IN BIOTECHNOLOGIES |
| Title of the subject | DRUG DESIGN AND DEVELOPMENT |
| Degree Course (class) | Industrial and Environmental Biotechnology (LM-8) |
| ECTS credits | 6 |
| Compulsory attendance | yes |
| Language | italian |
| Academic year | 2020-2021 |

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| Subject Teacher | |
| Name and Surname | Orazio Nicolotti |
| email address | orazio.nicolotti@uniba.it |
| Place and time of reception | by email |
| ECTS credits details | |
| | Discipline sector (SSD) |
| | CHIM/08 |
| | Area |
| | Chemistry |

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| Study plan schedule | Year of study plan | Semester | | |
| | II | I | | |
| Time management | | | | |
| | Lessons | Laboratory | Exercises | Total |
| CFU | 5 | 1 | | 6 |
| Total hours | 125 | 25 | | 150 |
| In-class study hours | 40 | 12 | | 52 |
| Out-of-class study hours | 85 | 13 | | 98 |

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| Syllabus | |
| Prerequisites / Requirements | |
| Expected learning outcomes (according to Dublin descriptors) | |
| Knowledge and understanding | Learning about main methods for the design of molecules of pharmacological or diagnostic interest. |
| Applying knowledge | Use of specific methodologies and technological platforms (molecular modeling) for the identification of molecular targets of biotechnological interest. |
| Making informed judgments and choices | Informed evaluation and interpretation of experimental data. |
| Communicating knowledge | Skills to analyze, propose and critically discuss the data with general audience. |

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| Capacities to continue learning | Critical approach to the medicinal chemistry topics. |
| Study Program | |
| Content | Fundamental interactions between molecules. Van der Waals forces. Hydrophobic effect. Ionic interactions. Dipolar interactions. 1D, 2D and 3D molecular descriptors. Quantitative Structure-Activity / Property Relationship. 1D, 2D and 3D molecular representation. Database searching. Similarity and molecular diversity. Principal component analysis, cluster analysis, genetic algorithms, multi-objective optimization techniques. Experimental design. Applicability domains. Basic of predictive toxicology. Pharmacophore models. QSAR models. Hansch equation. Free-Wilson equation. Craig Plot, CoMFA, GRID. Molecular mechanics. Conformational Analysis. Three-dimensional structure of proteins and active site. Protein Data Bank, Protein-Molecule Interaction, Molecular Docking. Molecular dynamics. De novo design. Virtual screening. |
| Bibliography and textbooks | <ul style="list-style-type: none"> • R. Leach, Molecular Modelling: Principles and Applications, Pearson Education EMA • Graham L. Patrick; Introduzione alla Chimica farmaceutica (EdiSES) • Gasco, Gualtieri, Melchiorre: Chimica Farmaceutica (Casa Editrice Ambrosiana) |
| Notes to textbooks | |
| Teaching methods | |
| Assessment methods (oral, written, ongoing assessment) | |
| Evaluation criteria (describe criteria for each of the above expected outcomes) | Knowledge of the basic methods for the design of molecules of pharmacological or diagnostic interest. Use of programs for molecular modeling. Critical evaluation of experimental results. |
| Further information | |