

<b>Module code</b>	SB-4342		
<b>Module Title</b>	Advanced Genetic Analysis		
<b>Degree/Diploma</b>	Bachelor of Science (Biology)		
<b>Type of Module</b>	Major Option		
<b>Modular Credits</b>	4	<b>Total student workload</b>	8 hours/week
		<b>Contact hours</b>	6 hours/week
<b>Prerequisite</b>	SB-2211 Genetics		
<b>Anti-requisite</b>	None		
<b>Aims</b>			
This module is designed for students to apply the combined power of classical and molecular genetics in investigating a wide range of biological questions. The students will also engage in discussion based on the primary literature to conceptualise a plan for studying key biological questions at the molecular level.			
<b>Learning Outcomes:</b>			
<i>On successful completion of this module, a student will be expected to be able to:</i>			
Lower order :	40%	<ul style="list-style-type: none"> <li>- Describe the definition of a gene</li> <li>- Describe gene organisation and chromosome structure</li> <li>- Describe the advantages and potential limitations of various model organism.</li> <li>- Understand the general principles of mutant analysis.</li> <li>- Identify mutation.</li> <li>- Discuss the various tools in reverse genetics.</li> <li>- Explain the concept and purpose of genome-wide mutant screens.</li> </ul>	
Middle order :	40%	<ul style="list-style-type: none"> <li>- Characterise the phenotype of a mutation.</li> <li>- Associate mutant phenotype to DNA sequence.</li> <li>- Classify mutation defects on the basis of its effect on the function.</li> <li>- Determine functionally related genes on the basis of their interaction and/or infer a logical order in which genes function in a pathway.</li> </ul>	
Higher order:	20%	<ul style="list-style-type: none"> <li>- Conduct a presentation on non-conventional model organisms used in research and discuss their limitations.</li> <li>- Work effectively in groups to develop a plan for investigating a key biological question.</li> </ul>	
<b>Module Contents</b>			
<ul style="list-style-type: none"> <li>- The basis for genetic analysis</li> <li>- Genomes, chromosomes and epigenetics</li> <li>- Model organisms and their genomes</li> <li>- Identifying and classifying mutants</li> <li>- Connecting phenotypes with DNA sequences</li> <li>- Mutant phenotypes and gene activity</li> <li>- Reverse genetics</li> <li>- Genome editing</li> <li>- Genome-wide mutant screens</li> <li>- Gene interactions: suppressors and enhancers</li> <li>- Epistasis and genetic pathways</li> </ul>			
<b>Assessment</b>	Formative assessment	Tutorials and feedback	
	Summative assessment	Examination: 60% Coursework: 40% <ul style="list-style-type: none"> <li>- Two (2) individual assignments (10%)</li> <li>- One (1) individual presentation (10%)</li> <li>- One (1) individual laboratory report (10%)</li> <li>- One (1) class test (10%)</li> </ul>	