

| Staff member | Organisation eg. SBS / IMS / PFR / LCR | Academic Group eg. EEB / BHB / CMPB / JGS | Size 60/90/120 eg. BScHons = 60pt, taught MSc = 60pt, BiomedHons = 90pt, MSc = 120pt | Title/Area | Expiry Date | Relevant Courses that students should take or have taken |
|---|--|---|--|--|-------------|---|
| Evolutionary genetics/genomics, Phylogeny projects | | | | | | |
| Nicholas Matzke | SBS | EEB | any | Phylogenetic biogeography - testing new models of diversification, extinction, dispersal on with clade of interest or simulated data | none | Evolution or biogeography courses such as: BIOSCI 109: Ecology & Evolution BIOSCI 210: Evolution - molecules to organisms BIOSCI 322: Genes, Populations, Species BIOSCI 700: Phylogenetics BIOSCI 395: Pacific Biogeography BIOSCI 731: Biogeography BIOSCI 220: Quantitative Biology |
| Nicholas Matzke | SBS | EEB | any | Structural phylogenetics - new models for including protein structure data in phylogenetics, using the models on protein groups of interest (e.g. bacterial flagellum), or testing models & performance on simulated or empirical data | none | Evolution courses such as: BIOSCI 109: Ecology & Evolution BIOSCI 210: Evolution - molecules to organisms BIOSCI 322: Genes, Populations, Species BIOSCI 700: Phylogenetics BIOSCI 220: Quantitative Biology |
| Nicholas Matzke | SBS | EEB | any | Structural phylogenetics - trialing methods of Multiple Sequence Alignment and MSA visualization for protein structure characters | | |
| Nicholas Matzke | SBS | EEB | any | Other potential phylogenetics/bioinformatics projects of interest, e.g. (1) make a bioinformatics tool (e.g. dot-plot visualiser for 3D protein structure characters), or (2) updating older questions with new data & methods: AMELX/AMELY (sex-specific proteins in mammal teeth), blood-clotting cascade, GULO pseudogene for vitamin C synthesis | | |
| Anna Santure | SBS | EEB | any | Recombination in hihi (stitchbird): why do males do it more? | none | BIOSCI 351 plus ideally BIOSCI 322 or BIOSCI 210; must take BIOSI 701 (okay if taking concurrently) |
| Anna Santure | SBS | EEB | any | Comparing mitochondrial and genomic signals of invasion in the common myna | none | BIOSCI 351 plus ideally BIOSCI 322 or BIOSCI 210; must take BIOSI 701 (okay if taking concurrently) |
| Nobuto Takeuchi | SBS | CMPB | any | Mathematical or computational modelling of genome evolution | none | BIOSCI 702, BIOSCI 700, BIOSCI 701 (700-level can be concurrent) |
| Nobuto Takeuchi | SBS | CMPB | Hons (60 or 90 pt), 120-pt MSc | Evolutionary genomics of prokaryote genomes | none | BIOSCI 700, BIOSCI 701, BIOSCI 702 (700-level can be concurrent) |
| Jane Allison | SBS | CMPB | any | Modelling how mitochondrial structure affects the flow of protons | | |
| Jane Allison | SBS | CMPB | any | Modelling antimicrobial peptide-membrane interactions | | |
| Sarah Knight | SBS | EEB | 90 or 120 | Divergence between NZ isolated yeast and international representatives | none | BIOSCI 220, ideally BIOSCI 322 and/or BIOSCI 351; BIOSCI 701 (700-level can be concurrent) |
| Sarah Knight | SBS | EEB | 120 | Population genetics of Wakame in NZ | none | BIOSCI 220, ideally BIOSCI 322 and/or BIOSCI 351; BIOSCI 701 (700-level can be concurrent) |
| Austen Ganley | SBS | BHB | any | Assessing AI models of genomic predictions | none | BIOSCI355 or equivalent |
| Austen Ganley | SBS | BHB | any | Determining functional areas of genomes using random DNA | none | BIOSCI355 or equivalent |
| Austen Ganley | SBS | BHB | 90 or 120 | Determining the dominant recombination pathway in the ribosomal RNA gene repeats | none | BIOSCI351 or BIOSCI355 or equivalent; BIOSCI736 (700-level can be concurrent) |
| Austen Ganley | SBS | BHB | 90 or 120 | Experimental evolution of random DNA sequences | none | BIOSCI210 or BIOSCI355 or equivalent |
| Kim Handley | SBS | EEB | 120 | Determine CRISPR spacer variation within bacterial populations as a dynamic record of bacterial-viral interactions | none | BIOSCI 701 (700-level can be concurrent), ideally BIOSCI 204 and 347 |
| Simon Greenhill | SBS | EEB | 90 or 120 | Tracing the evolution of the Polynesian gods | none | Evolution courses such as: BIOSCI 109: Ecology & Evolution, BIOSCI 210: Evolution - molecules to organisms, BIOSCI 322: Genes, Populations, Species, BIOSCI 700: Phylogenetics, BIOSCI 220: Quantitative Biology |
| Simon Greenhill | SBS | EEB | 90 or 120 | Testing and evaluating models of language evolution | none | BIOSCI700 |
| Simon Greenhill | SBS | EEB | 90 or 120 | Using phylogenies of language to shed light into human prehistory | none | Evolution courses such as: BIOSCI 109: Ecology & Evolution, BIOSCI 210: Evolution - molecules to organisms, BIOSCI 322: Genes, Populations, Species, BIOSCI 700: Phylogenetics, BIOSCI 220: Quantitative Biology |
| Priscila Salloum | SBS | EEB | 120 | Exploring the genetic basis of shell colour in chitons | | 2029 BIOSCI 355 or equivalent |
| Marine/Freshwater Projects | | | | | | |
| Kim Handley | SBS | EEB | 120 | Determine the function of hellorhodopsin in the photoheterotrophy of estuarine bacteria | none | BIOSCI 204, ideally BIOSCI 347 |
| Kim Handley | SBS | EEB | 120 | Confirm predictions of anaerobic nitrogen metabolism in the subsurface Nitrospirota bacterial phylum | none | BIOSCI 204, ideally BIOSCI 347 |
| Brendon Dunphy | SBS | EEB | any | Ecophysiology of seagrids | none | BIOSCI 334 |
| Kendall Clements | SBS | CMPB | 120 | Diet in herbivorous and detritivorous coral reef fishes | none | BIOSCI 335, BIOSCI 725 |
| Kendall Clements | SBS | CMPB | 120 | Biodiversity of coral reef cyanobacteria | | |
| Priscila Salloum | SBS | EEB | 120 | Probiotics and experimental microbiome manipulation in New Zealand chitons | | 2029 BIOSCI 204, BIOSCI 749 can be helpful but not required |
| Zoology/Behaviour/Biodiversity Projects | | | | | | |
| Sarah Knight | SBS | EEB | 120 | Metabolic potential of Aotearoa NZ's yeasts | none | Ideally BIOSCI 322, 348, 204 but not necessary |
| Sarah Knight | SBS | EEB | 120 | Biodiversity of NZ vineyards | none | BIOSCI 220, ideally BIOSCI 322 and/or BIOSCI 351; BIOSCI 701 (700-level can be concurrent). This is a computational project with existing data. |
| Manpreet Dhami | SBS | EEB | 120 | Microbiomes of endemic pollinators | none | skills: microbial ecology, microbiology, entomology, bioinformatics, statistics |
| Manpreet Dhami | SBS | EEB | 120 | Microbiome of kiwi | none | skills: microbial ecology, microbiology, conservation biology, bioinformatics statistics |
| Manpreet Dhami | SBS | EEB | 120 | Competition coexistence dynamics of nectar yeast communities | none | skills: ecology, stats, microbiology |
| Libby Liggins | SBS | EEB | 120 | Monitoring of tropical and subtropical reef fishes using citizen science | none | |
| Libby Liggins | SBS | EEB | 60/120 | Using metabarcoding of Autonomous Reef Monitoring Systems to evaluate the shallow marine biodiversity of Rangitāhua | none | |
| Priscila Salloum | SBS | EEB | any | Microbiome profiling in chitons | | 2029 BIOSCI 204 may be helpful, but not required |
| Priscila Salloum | SBS | EEB | any | Shell colour variation, colour quantification and camouflage in chitons | | 2029 BIOSCI 109 |
| Neil Birrell & Rich Leschen | SBS/MWL | EEB | 120 | Finding Mates in Flightless Beetles: Comparative morphology of cerambycid antennal ultrastructures | none | BIOSCI 208 and 338 wpu'd be helpful but not necessary |
| Plant and Other Terrestrial Projects | | | | | | |
| Margaret Stanley | SBS | EEB | 120 | Non-chemical weed management - using native pathogens. Māori student preferred for funding (Bioprotection Aotearoa) - fees + \$10K stipend funding | | BIOSCI 394 ideally, plant pathology |
| Cate Macinnis-Ng | SBS | EEB | 60, 120 | Plant responses to a changing climate - field and glasshouse-based projects available | none | BIOSCI 325, BIOSCI 766 |
| James Brock | SBS | EEB | 120 | Ferns and fire - spore banks in a changing landscape (must start July, have driving licence) | none | BIOSCI 394 and maybe 325 |
| Chris Carrie | SBS | CMP | 60/120 | Determining the molecular basis of the high temperature stress resistance of geothermal Kānuka | none | BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochemistry, genetics or molecular biology |

| Project Overview | | | | | | |
|---|-----------|----------------|-----------|---|--|--|
| Project Title | | Project Leader | | Project Description | | |
| Biomedical and Human Biology | | | | | | |
| Inken Kelch | SBS | BHB | 90 or 120 | Comprehensive multicolour imaging of immune organs | none | immunology background (MEDSCI 202 or BIOSCI 201, ideally MEDSCI 314) |
| Alicia Didsbury & Daniel Verdon | SBS | BHB | 90 or 120 | Next-Generation T-Cell Expansion for Adoptive Immunotherapy: Integrating Serum-Free Media and Novel Culture Platforms | none | immunology background (MEDSCI 202 or BIOSCI 201, ideally MEDSCI 314) |
| Catherine (Kate) Angel | SBS | BHB | 60 | Analysis of RNA-seq data to determine how tPA treatment impairs the brain endothelial barrier following ischemic stroke | Earliest Start: BIOSCI 761 2nd Sem 2026 | BIOSCI 220 & stage III/700 level courses with cellular & molecular focus. Experience with R packages. |
| Christopher Walker | SBS | BHB | | Cell signalling and receptors in metabolic disease | | |
| Hilary Sheppard | SBS | BHB | 90 or 120 | Gene editing for severe skin conditions | | |
| Jessie Jacobsen | SBS | BHB | | Lymphatic modulation by | | |
| John Taylor | SBS | BHB | | | | |
| Anthony Phillips | SBS | BHB | 120 | Oxidative stress measures at the bedside - developing a new window on critical illness | | do 761 in S1 |
| Anthony Phillips | SBS | BHB | 120 | Extending the biological capability of a lowgravity simulation machine | | do 761 in S1; interest in engineering and some basic programming experience advantageous |
| Anthony Phillips | SBS | BHB | 120 | Modulating lymphatic function to help congestive heart failure | | do 761 in S1; interest in engineering and some basic programming experience advantageous |
| Anthony Phillips | SBS | BHB | 120 | Attenuating negative effects of space flight on mitochondria | | do 761 in S1 |
| Garth Cooper | SBS | BHB | | | | |
| Kerry Loomes | SBS | BHB | | | | |
| Mike Taylor | SBS | BHB | | | | |
| Rod Dunbar | SBS | BHB | | | | |
| Russell Snell | SBS | BHB | | | | |
| Saem Park | SBS | BHB | 90 or 120 | Spatial Mapping of Immune Responses in Human Cancers | | immunology background (MEDSCI 202 or BIOSCI 201, ideally MEDSCI 314) |
| Nicole Edwards | SBS | BHB | | | | |
| Emma Scotter | SBS | BHB | 90 or 120 | Developing cell models of neurodegenerative disease | | |
| Emma Scotter | SBS | BHB | 90 or 120 | Multiplex immunohistochemistry of human brain tissue in motor neuron disease | | |
| Jennifer Miles-Chan | SBS | BHB | | | | |
| Paul Harris | SBS | BHB | | | | |
| Shaun Lott | SBS | CMPB/BHB | 60/90/120 | RNaseHi as a target for new antibiotics | | |
| Iman Kavianinia | SBS | | 120 | Developing Next-Generation Platforms for Precision Cancer Therapy | | |
| Cellular, Molecular and Physiological Biology | | | | | | |
| Matthew Sullivan | SBS | CMP | 120 | Structural and Functional Analysis of Protein-Metallodrug Interactions | none | BIOSCI350, BIOSCI353 or equivalent. |
| David Goldstone | SBS | CMP | 120 | Using structural biology to understand the function and role of polygalacturonases in fruit ripening | none | Biochemistry or plant biology. |
| David Goldstone | SBS | CMP | 121 | Structural investigation of anti-retroviral compounds targeting the HIV-1 Capsid | none | Biochemistry, BIOSCI350, BIOSCI353 or equivalent. |
| David Goldstone | SBS | CMP | 120 | Establishing a method to probe the recognition of the retroviral capsids by the host restriction factor Trim5alpha. | none | BIOSCI350, BIOSCI353 or equivalent. |
| Karine David | SBS | CMP | 120 | Molecular pathways controlling fruit flesh development in apple/tomato | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Karine David | SBS | CMP | 120 | Analysis on candidate genes controlling fruit size in kiwifruit | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Soledad Perez Santangelo | SBS | CMP | 60/120 | Role of alternative splicing in circadian clock regulation by temperature in legumes | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Soledad Perez Santangelo | SBS | CMP | 60/120 | Analysis of candidate genes involved in temperature-mediated modulation of circadian rhythms in legumes | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Chris Carrie | SBS | CMP | 60/120 | Determining the molecular basis of the high temperature stress resistance of geothermal <i>Kanuka</i> | none | BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochemistry, genetics or molecular biology |
| Chris Carrie | SBS | CMP | 60/120 | What is the molecular role of the MICOS complex in maintaining plant mitochondrial ultrastructure | none | BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochemistry, genetics or molecular biology |
| Chris Carrie | SBS | CMP | 60/120 | Development of an inducible chloroplast biogenesis system | none | BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochemistry, genetics or molecular biology |
| Kim Snowden | PFR | CMP | 60/120 | Understanding the pathways that control plant architecture | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Kim Snowden | PFR | CMP | 60/120 | Developing new tools to control plant development | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Kim Snowden | PFR | CMP | 60/120 | Gene editing of crop plants to improve management and yield | none | BIOSCI 326, BIOSCI 351 or equivalent |
| Augusto Simoes-Barbosa | SBS | CMP | | | | |
| Iain Hay | SBS | CMP | 120 | Designing protein inhibitors for bacterial outer membrane proteins | none | |
| Iain Hay | SBS | CMP | 120 | Understanding bacterial intra species competition and kin recognition | none | |
| Richard Kingston | SBS | CMP | 120 | How does Syto9, a widely-used fluorescent dye, bind nucleic acids? | | |
| Richard Kingston | SBS | CMP | 120 | Investigating the oligomeric state of the coiled-coil found in Paramyxoviral Phosphoproteins. | | |
| Andrew Allan | SBS | CMP | | | | |
| Anthony Poole | SBS | CMP | 120 | Recovery and sequencing of complete mitogenomes using Oxford nanopore technology | | |
| Craig Millar | SBS | CMP | 120 | Establishing parentage and relatedness in the communal groups of Brown Skua using microsatellite markers | | |
| Matthew Templeton | SBS | CMP | | | | |
| Christopher Squire | SBS | CMP | 120 | How immune receptors switch on: exploring ligand binding with molecular dynamics simulations | | |
| Christopher Squire | SBS | CMP | 120 | Predicting resistance mutations: using molecular dynamics to understand how rodenticides fail | | |
| Ghader Bashiri | SBS | CMP | 120 | Genome mining for novel antimicrobial agents | | |
| Ghader Bashiri | SBS | CMP | 120 | Biosynthetic enzymes in microbial secondary metabolism | | |
| Ghader Bashiri | SBS | CMP | 120 | Enzyme engineering for biocatalysis | | |
| Ghader Bashiri | SBS | CMP | 120 | How pathogenic bacteria sense oxidative stress? | | |
| Ghader Bashiri | SBS | CMP | 120 | Targeting DNA repair for new antibiotics | | |
| Robert Schaffer | SBS | CMP/PFR | | | | |
| Robin MacDiarmid | SBS | CMP/PFR | | | | |
| Shaun Lott | SBS | CMPB/BHB | 60/90/120 | Rhs repeat proteins in bacterial competition and multicellular evolution. | | |
| Tristan de Rond | SCS | CMP | 60/90/120 | Genome mining / biosynthetic enzyme discovery | | Biochemistry, Molecular biology, Organic chemistry |
| Tristan de Rond | SCS | CMP | 60/90/120 | Enzyme biotechnology / synthetic biology / metabolic engineering | | Biochemistry, Molecular biology, Organic chemistry. Some computer competency would be good if you want to use the pipetting robot :) |
| Tristan de Rond | SCS | CMP | 60/90/120 | Bioinformatic tool development for genome mining / biosynthetic enzyme discovery | | Microbial genomics, Bioinformatics, Biochemistry |
| Irina Miller | Daisy Lab | | 60 | Investigating Flocculation in Genetically Engineered <i>Pichia pastoris</i> for Heterologous Protein Expression - how flocculation can be used to reduce downstream processing in heterologous protein expression. 1 to 2 projects will be offered in this topic. | for BIOSCI 761 in S1 2026. Research completed end of S2 2026 | Background in Protein Structure/Function, microbiology |
| Irina Miller | Daisy Lab | | 60 | Computational analysis of fermentation data for protein optimisation in heterologous expression systems. 1 to 2 projects will be offered in this topic. | for BIOSCI 761 in S1 2026. Research completed end of S2 2026 | Background in Protein Structure/Function, microbiology |
| Irina Miller | Daisy Lab | | 60 | Investigation of the Relationship Between Glycosylation Patterns and Functional Properties of Recombinant Lactoferrin. 1 to 2 projects will be offered in this topic. | for BIOSCI 761 in S1 2026. Research completed end of S2 2026 | Background in Protein Structure/Function, microbiology |
| Irina Miller | Daisy Lab | | 60 | Stability assessment of Recombinant Lactoferrin under variable processing and storage conditions. 1 to 2 projects will be offered in this topic. | for BIOSCI 761 in S1 2026. Research completed end of S2 2026 | Background in Protein Structure/Function, microbiology |
| Irina Miller | Daisy Lab | | 60 | Functional testing of recombinant lactoferrin. 1 to 2 projects will be offered in this topic. | for BIOSCI 761 in S1 2026. Research completed end of S2 2026 | Background in Protein Structure/Function, microbiology |
| Projects at the policy interface | | | | | | |

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|------------------------------|-----------------|-----|---------|---|---------------------------|---|
| Juliet Gerrard | SBS/SCS/CHEMMAT | any | any | Bespoke projects to suit student interest at the policy interface | none | any |
| Ecology and Evolution | | | | | | |
| David Pattemore | MWL | EEB | | | | |
| Mai Padamsee | MWL | EEB | | | | |
| Shane Wright | SBS | EEB | | | | |
| Alexei Drummond | SBS | EEB | | | | |
| Cate Macinnis-Ng | SBS | EEB | | | | |
| Gavin Lear | SBS | EEB | any | Microbial interactions with plastic pollutants | | |
| Greg Holwell | SBS | EEB | any | Camouflage, aposematism, mimicry, and other antipredator adaptations in insects (variety of possible projects) | none | BIOSCI 207, 337,338 |
| Jacqueline Beggs | SBS | EEB | | | | |
| James Russell | SBS | EEB | | | | |
| Maren Wellenreuther | PFR | EEB | | | | |
| Mary Sewell | SBS | EEB | 120 | Lipid analysis of the roe, eggs and embryos of the sea urchin <i>Centrostephanus rodgersii</i> | none | BIOSCI 208 or knowledge of invertebrates useful. Good background in chemistry. |
| Rochelle Constantine | SBS | EEB | | | | |
| Tony Hickey | SBS | EEB | | | | |
| ZhiQiang Zhang | MWL | EEB | | | | |
| Dave Seldon & Sam Brown | SBS | EEB | 90, 120 | can micro-hymenoptera diversity indicate arthropod diversity in mangroves | | |
| Dave Seldon | SBS | EEB | 120 | Revision of aotearoa weevil genus <i>Clypeolus</i> | | |
| Dave Seldon | SBS | EEB | 90,120 | micro-hymenoptera diversity in urban restoration projects | | |
| Dave Seldon | SBS | EEB | 120 | Revision of the infinite species group within the genus <i>Mecodema</i> (ground beetles) | | |
| Al Glen | SBS | EEB | 120 | Revision of the endemic ground beetle genus <i>Neoferonia</i> | | |
| Anne Gaskett | SBS | EEB | any | Sensory ecology and interactions, esp. colour and light perception (plant-animal or fungi-animal interactions such as pollination or fruit dispersal, or seabirds and sensory traps such as plastic ingestion or light pollution) | none | any zoology, botany and natural history subjects (e.g. BIOSCI 206, BIOSCI 207, BIOSCI325, BIOSCI337, BIOSCI 338, BIOSCI395) |
| Brendon Dunphy | SBS | EEB | 120 | Physiological approaches to improve invertebrate aquaculture | none | BIOSCI 208, MARSCI328, and BIOSCI 727 |
| Bruce Burns | SBS | EEB | | | | |
| Darren Ward | MWL | EEB | any | Do citizen scientists (Naturalist) and museum collections document similar data on biodiversity | none | |
| Darren Ward | MWL | EEB | any | Using automated identification models to identify invasive ants (can include a small field component) | none | |
| Emma Carroll | SBS | EEB | | | | |
| J David Aguirre | SBS | EEB | 120 | Heritability of tropical damselfish feeding and locomotion traits | none | |
| J David Aguirre | SBS | EEB | 120 | Diet niche variance and segregation within populations of damselfishes using Stable Isotope analyses | none | |
| Kristal Cain | SBS | EEB | | | | |
| Libby Liggins | SBS | EEB | 120 | Recent and predicted future changes to NZ shallow marine environments using dispersal simulations and niche modelling | none | |
| Libby Liggins | SBS | EEB | 120 | Diet analysis of co-occurring NZ urchins using metabarcoding | none | |
| Libby Liggins | SBS | EEB | 120 | Kin and spatial genetic structuring of tropical damselfishes across reefscape | none | |
| Margaret Stanley | SBS | EEB | 120pt | Bioprotection Aotearoa funded project on weed management (w stipend) for Māori student | for BIOSCI 761 in S2 2026 | |
| Alice Della Penna | IMS+SBS | EEB | Any | Investigating plankton patchiness with semi-automated microscopy | | |