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		Academic				
	Organisation eg.		/ Size 60/90/120 eg. BScHons =			Relevant Courses that students should take or have taken
	SBS / IMS / PFR /	BHB / CMPB /				
Staff member	LCR	JGS	BiomedHons = 90pt, MSc = 120pt	Title/Area	Expiry Date	
Evolutionary						
genetics/genomics, Phylogeny						
projects						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
				Phylogenetic biogeography - testing new models of diversification, extinction, dispersal on with clade of interest or		BIOSCI 731: Biogeography
Nicholas Matzke	SBS	EEB	anv	simulated data	none	BIOSCI 220: Quantitative Biology
						Evolution courses such as:
						BIOSCI 109: Ecology & Evolution
						BIOSCI 210: Evolution - molecules to organisms
						BIOSCI 322: Genes, Populations, Species
						BIOSCI 700: Phylogenetics
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	BIOSCI 220: Quantitative Biology
INICIOIDS IVIDIZACE	203	LED	arry	protein groups of interest (e.g. bacterial flagellum), or testing models & performance on simulated or empirical data Structural phylogenetics - trialing methods of Multiple Sequence Alignment and MSA visualization for protein structure	none	DIOSCI 220. Quantitative Diology
Nicholas Matzke	SBS	EEB	any	characters		
			<u>'</u>	Other potential phylogenetics/bioinformatics projects of interest, e.g. (1) make a bioinformatics tool (e.g. dot-plot		
				visualiser for 3Di 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		
Nicholas Matzke	SBS	EEB FFB	any	synthesis		
Anna Santure Anna Santure	SBS SBS	EEB	any	Recombination in hihi (stitchbird): why do males do it more? 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) 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	СМРВ	any	Modelling how mitochondrial structure affects the flow of protons		
Jane Allison	SBS	СМРВ	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)
	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	внв	any	Assessing Al models of genomic predictions	none	BIOSCI355 or equivalent
Austen Ganley Austen Ganley	SBS SBS	BHB	any 90 or 120	Determining functional areas of genomes using random DNA Determining the dominant recombination pathway in the ribosomal RNA gene repeats	none	BIOSCI355 or equivalent BIOSCI351 or BIOSCI355 or equivalent; BIOSCI736 (700-level can be concurrent)
Austen Ganley 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
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						Evolution courses such as: BIOSCI 109: Ecology & Evolution, BIOSCI 210: Evolution - molecules to organisms,
Simon Greenhill	SBS	EEB	90 or 120	Tracing the evolution of the Polynesian gods	none	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
						Evolution courses such as: BIOSCI 109: Ecology & Evolution, BIOSCI 210: Evolution - molecules to organisms,
Simon Greenhill	SBS	EEB	90 or 120	Using phylogenies of language to shed light into human prehistory	none	BIOSCI 322: Genes, Populations, Species, BIOSCI 700: Phylogenetics, BIOSCI 220: Quantitative Biology
Marine/Freshwater Projects						, , , , , , , , , , , , , , , , , , , ,
Kim Handley	SBS	EEB	120	Determine the function of heliorhodopsin 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 seabirds	none	BIOSCI 334
Kendall Clements	SBS	СМРВ	120	Diet in herbivorous and detritivorous coral reef fishes	none	BIOSCI 335, BIOSCI 725
Kendall Clements	SBS	СМРВ	120	Biodiversity of coral reef cyanobacteria		
Zoology/Behaviour/Biodiversit						
y Projects Sarah Knight	SBS	EEB	120	Metabolic potential of Aotearoa NZ's yeasts	none	Ideally BIOSCI 322, 348, 204 but not necessary
Sur un miligint	303		110	metabone potential of Motedial NE 3 years	none.	BIOSCI 220, ideally BIOSCI 322 and/or BIOSCI 351; BIOSCI 701 (700-level can be concurrent). This is a
Sarah Knight	SBS	EEB	120	Biodiversity of NZ vineyards	none	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	FFB	60/120	Using metabarcoding of Autonomous Reef Monitoring Systems to evaulate the shallow marine biodiversity of Rangitāhua	none	
	SBS/MWL	EEB	120	Finding Mates in Flightless Beetles: Comparative morphology of cerambycid antennal ultrastructures	none	BIOSCI 208 and 338 wpuld be helpful but not necessary
Plant and Other Terrestial	,			gg. ngg. ngg		
Projects						
				Non-chemical weed management - using native pathogens. Māori student preferred for funding (Bioprotection		
Margaret Stanley	SBS	EEB	120	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 Chris Carrie	SBS	CMP	120 60/120	Ferns and fire - spore banks in a changing landscape (must start July, have driving licence) Determining the molecular basis of the high temperature stress resistance of geothermal Kānuka	none	BIOSCI 394 and maybe 325 BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochmestry, genetics or molecular biology
Cinis carrie	303	CIVIE	00/120	Determining the morecular basis of the high temperature stress resistance of geothernial Kaffuka	none	503Ci 320, 5103Ci 332 oi equivalent. Anyone interested in biochinestry, genetics of molecular biology
Biomedical and Human Biology						
Inken Kelch	SBS	внв	90 or 120	Comprehensive multicolour imaging of immune organs	none	immunology background (MEDSCI 202 or BIOSCI 201, ideally MEDSCI 314)

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Alicia Didsbury & Daniel Verdon	SBS	внв	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)
Cathorino (Kato) Angol	cnc	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.
Catherine (Kate) Angel Christopher Walker	SBS	ВНВ	80	Cell signalling and receptors in metabolic disease	Earliest Start. BIOSCI 701 2110 3em 2020	BIOSCI 220 & Stage III/700 level courses with Centular & Molecular Tocus, Experience with K packages.
	SBS	внв	90 or 120	Gene editing for severe skin conditions		
Jessie Jacobsen	SBS	внв		Lymphatic modulation by		
John Taylor	SBS	ВНВ				
	SBS	BHB	120	Oxidative stress measures at the bedside - developing a new window on critical illness		do 761 in S1
Anthony Phillips Anthony Phillips	SBS	ВНВ	120	Extending the biological capability of a lowgravity simulation machine Modulating lymphatic function to help congestive heart failure		do 761 in S1; interest in engineering and some basic programming experience advantageous do 761 in S1; interest in engineering and some basic programming experience advantageous
	SBS	ВНВ	120	Attenuating negative effects of space flight on mitochondria		do 761 in S1
Garth Cooper	SBS	внв				
Kerry Loomes	SBS	ВНВ				
	SBS	ВНВ				
Rod Dunbar	SBS	BHB				
Russell Snell 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	50 01 120	Spacial Wapping of Milliane Responses in Human Cancers		immunology background (PIEDSOT 202 of BIOSOT 201, Ideally PIEDSOT 314)
	SBS	ВНВ	90 or 120	Developing cell models of neurodegenerative disease		
	SBS	ВНВ	90 or 120	Multiplex immunohistochemistry of human brain tissue in motor neuron disease		
Jennifer Miles-Chan	SBS	ВНВ				
Paul Harris	SBS	BHB	50 (00 (4 00	Pale at the second for a second		
Shaun Lott	SBS	CMPB/BHB	60/90/120	RNaseHI as a target for new antibiotics	1	
Iman Kavianinia Cellular, Molecular and	SBS		120	Developing Next-Generation Platforms for Precision Cancer Therapy		
Physiological Biology						
	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
	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 Chris Carrie	SBS SBS	CMP	60/120 60/120	Analysis of candidate genes involved in temperature-mediated modulation of circadian rhythms in legumes Determining the molecular basis of the high temperature stress resistance of geothermal Kānuka	none	BIOSCI 326, BIOSCI 351 or equivalent BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochmestry, 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 Biochmestry, genetics or molecular biology
	SBS	CMP	60/120	Development of an inducible chloroplast biogenesis system	none	BIOSCI 326, BIOSCI 351 or equivalent. Anyone interested in Biochmestry, 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
KIIII SIIOWACII	PFR	CMP	60/120	Gene editing of crop plants to improve management and yield	none	BIOSCI 326, BIOSCI 351 or equivalent
Augusto Simoes-Barbosa Iain Hay	SBS SBS	CMP	120	Designing protein inhibitors for bacterial outer membrane proteins	none	
	SBS	CMP	120	Understanding bacterial intra species competion 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.		
	SBS	CMP				
	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 Christopher Squire	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	1	
Robert Schaffer Robin MacDiarmid	SBS	CMP/PFR CMP/PFR				
Shaun Lott	SBS	CMP/PFK CMPB/BHB	60/90/120	Rhs repeat proteins in bacterial competition and multicellular evolution.	1	
	SCS	CMP	60/90/120	Genome mining / biosynthetic enzyme discovery		Biochemistry, Molecular biology, Organic chemistry
			· ·			Biochemistry, Molecular biology, Organic chemistry. Some computer competency would be good if you want
Tristan de Rond	SCS	CMP	60/90/120	Enzyme biotechnology / synthetic biology / metabolic engineering		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
				Investigating Flocculation in Genetically Engineered Pichia Pastoris for Heterologous Protein Expression - how flocculation can be used to reduce downstream processing in heterologous protein expression. 1 to 2 projects will be	for BIOSCI 761 in S1 2026. Research completed	
Irina Miller	Daisy Lab		60	offered in this topic.	end of S2 2026	Background in Protein Structure/Function, microbiology
				Computational analysis of fermentation data for protein optimisation in heterologous expression systems. 1 to 2	for BIOSCI 761 in S1 2026. Research completed	g
Irina Miller	Daisy Lab		60	projects will be offered in this topic.	end of S2 2026	Background in Protein Structure/Function, microbiology
				Investigation of the Relationship Between Glycosylation Patterns and Functional Properties of Recombinant Lactoferrin		
Irina Miller	Daisy Lab		60	1 to 2 projects will be offered in this topic.	end of S2 2026	Background in Protein Structure/Function, microbiology
Inima Malillan	Dairestak		60	Stability assessment of Recombinant Lactoferrin under variable processing and storage conditions. 1 to 2 projects will	for BIOSCI 761 in S1 2026. Research completed	Designation of the Company of Com
Irina Miller	Daisy Lab		bU	be offered in this topic.	end of S2 2026 for BIOSCI 761 in S1 2026. Research completed	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.	end of S2 2026	Background in Protein Structure/Function, microbiology
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Projects at the policy interface						
Juliet Gerrard	SBS/SCS/CHEMMAT	any	any	Bespoke projects to suit student interest at the policy interface	none	any
Ecology and Evolution						

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David Pattemore	MWL	EEB				
Maj 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	I	
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 Centrostephanus rodgersii	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 Clypeolus		
Dave Seldon	SBS	EEB	90,120	micro-hymenotpera diversity in urban restoration projects		
Dave Seldon	SBS	EEB	120	Revision of the infimate species group within the genus Mecodema (ground beetles)		
Al Glen	SBS	EEB	120	Revision of the endemic ground beetle genus Neoferonia		
				Sensory ecology and interactions, esp. colour and light perception (plant-animal or fungi-animal interactions such as		any zoology, botany and natural history subjects (e.g. BIOSCI 206, BIOSCI 207, BIOSCI325, BIOSCI337, BIOSCI
Anne Gaskett	SBS	EEB	anv	pollination or fruit dispersal, or seabirds and sensory traps such as plastic ingestion or light pollution)	none	338. BIOSCI395)
Brendon Dunphy	SBS	EEB	120	Physiological apporaches to improve invertebrate aquaculture	none	BIOSCI 208. MARSCI328, and BIOSCI 727
Bruce Burns	SBS	EEB		, , , , , , , , , , , , , , , , , , , ,		
Darren Ward	MWL	EEB	anv	Do citizen scientists (iNaturalist) and museum collections document similar data on biodiversity	none	
Darren Ward	MWL	EEB	anv	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 segragation within populations of damselfishes using Stable Isotope analyses	none	
Kristal Cain	SBS	EEB				
				Recent and predicted furture changes to NZ shallow marine environments using dispersal simulations and niche		
Libby Liggins	SBS	EEB	120	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 reefscapes	none	
LIDDY LIGGIIIS	363	CCD	120	All and spatial genetic structuring of tropical damsemsiles across reciscapes	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	Anv	Investigating plankton patchiness with semi-automated microscopy		
Auto Deno i elilla			ruiy	mestigating painton pateriness with semi-activated microscopy		