2026/2027

Univerza v Mariboru
Medicinska fakulteta

UČNI NAČRT PREDMETA / COURSE SYLLABUS	
Ime predmeta:	**Molekularna biologija z genetiko**
Course title:	**Molecular biology with genetics**

Študijski program in stopnja Study programme and cycle	Študijska smer Study option	Letnik Year of study	Semester Semester
Splošna medicina, enovit magistrski študijski program		Prvi	2.
General medicine, Uniform master's degree study program		First	2nd

Vrsta predmeta (obvezni ali izbirni) / Course type (compulsory or elective)	obvezni compulsory
Univerzitetna koda predmeta / University course code:	

Predavanja Lectures	Seminar Seminar	Vaje Tutorial			Klinične vaje Clinical training	Druge oblike študija Other forms of study	Samost. delo Individual work	ECTS
		AV	LV	RV				
30	15		15				90	5

Nosilec predmeta / Course coordinator:	**doc. dr. Špela Stangler Herodež**

Jeziki /Languages:	**Predavanja / Lectures:**	slovenski/slovene
	Vaje / Tutorial:	slovenski/slovene

Pogoji za vključitev v delo oz. za opravljanje študijskih obveznosti:

ŠTUDIJSKE OBVEZNOSTI ŠTUDENTOV

30 ur predavanj je razdeljenih med štiri predavatelje, ki samostojno pripravijo izpitna vprašanja na svoje teme. Udeležba na predavanjih je zelo priporočljiva, ker se predavanja, seminarji in vaje prepletajo in dopolnjujejo.

15 ur vaj je razporejenih v 5 sklopov:

• Vsi študenti si morajo preskrbeti Navodila za vaje, ki jih lahko kupijo ali si sposodijo v knjižnicah.

• Študenti se morajo pred pričetkom vaj seznaniti z navodili za opravljanje vaj in varno delo (nevarnosti pri delu, delo s kemikalijami in infektivnim materialom), ki so napisana na koncu Navodil za vaje.

Prerequisites for enrolling in the course or for performing study obligations:

ACADEMIC OBLIGATIONS OF STUDENTS

30 hours of lectures are divided between four lecturers who independently prepare examination questions on their topics. Attendance at lectures is highly recommended because the lectures, seminars and tutorials intertwined and are complementary.

15 hours of laboratory work is arranged in 5 sets.

• All students must provide instructions for exercises that can be bought or borrowed in the library of Medical faculty.
It is necessary to be prepared in advance for the course.

• Students should be aware of before starting the courses with instructions for performing exercises and work safely (hazard at work, working with chemicals and infective material), which are written at the end of the Instructions

15 ur seminarjev se opravlja v treh skupinah, pri štirih predavateljih. Vsak predavatelj pripravi seminarske teme, ki so objavljene v datumskem razporedu. Študenti se predhodno pripravijo na seminar tako, da preštudirajo snov po zapiskih in ustrezna navedena poglavja po priloženi literaturi. Predavatelj izvede krajše uvodno predavanje, ki mu sledijo naloge in diskusija. Študenti rešujejo računske in problemske naloge v času seminarjev ali dobijo naloge na predavanjih in jih individualno ali po dva skupaj rešijo do seminarske ure, kjer se naloge predstavi in o njih diskutira. Študenti oddajo v pisni obliki poročilo, povzetek diskusije oz. rešene problemske naloge. Teme seminarjev so zajete tudi pri pisnem izpitu, zato je poznavanje vsebin seminarskih tem nujno za dobro opravljen izpit. Prisotnost na seminarjih je obvezna in se preverja. POGOJI ZA PRISTOP K POSAMEZNEMU PREVERJANJU ZNANJA: (Opravljene naloge pri vajah in seminarjih so pogoj za pristop k pisnemu izpitu.) • Uspešno opravljene vaje so pogoj za pristop k izpitu.	for laboratory work. 15 hours of seminars are carried out in three groups with four professors. Each lecturer will prepare seminar topics, which are published in the schedule date range. Students are conditioned to the seminar so that they study the notes of the lectures and the relevant chapters of the mentioned accompanying literature. The lecturer conducted a short introductory lecture, followed by tasks and discussion. Students solve computational tasks and problem during seminars or receive tasks in class and individual or two together to solve seminar problems. Students submitting in writing a report, or a summary of the discussion. problem solved tasks. Topics seminars are also covered in the exam, so mastering the contents of this seminar is essential for good exam. Attendance at seminars is mandatory and shall be checked. REQUIREMENTS FOR ACCESS TO INDIVIDUAL KNOWLEDGE CHECKING: (Laboratory and seminar work project is condition for approaching to written exam.) • Completion of assignments are a prerequisite for the written exam.

Vsebina (kratek pregled učnega načrta):	**Content (syllabus outline):**
1.Vloga molekularne biologije in genetike v medicini.	1. The role of molecular biology and genetics in medicine.
2. Struktura, morfologija in klasifikacija humanih kromosomov.	2. Structure, morphology and classification of human chromosomes.
3. Osnove citogenetike, molekularna citogenetika, klinična citogenetika – sindromologija.	3. The principles of cytogenetics, molecular cytogenetics, clinical cytogenetics (syndromology).
4. Spolni kormosomi, X inaktivacija napake spolnih kromosomov.	4. Sex chromosomes, X inactivation, sex chromosomes aberrations.
5. Dedovanje, Mendelejevi zakoni, dominantno, recesivno dedovanje, atipični vzorci dedovanja.	5. Inheritance, Mendelien laws, dominant, recessive single-gene inheritance, atypical patterns of inheritance.
6. Struktura in lastnosti DNA molekule.	6. Structure and features of DNA molecule.
7. Podvajanje DNA: razlike med prokarionti in evkarionti.	7. DNA replication: prokaryotes, eukaryotes.
8. DNA mutacije, popravljanje DNA napak.	8. DNA mutations, repairing mechanisms.
9. RNA, transkripcija, translacija, vrste RNA, ribocimi.	9. RNA, transcription, translation, ribocimes
10. Procesiranje proteinov, posttranskripcijske modifikacije proteinov.	10. Protein processing, posttranslational modifications of proteins.
11. Človeški genom, jederni genom, genetski kod,	11. Human genome, nuclear genome, genetic code,

zgradba gena, genske družine, tandemske ponovitve, mitohondrijski genom.	structure of the gene, gene families, tandem repeats, mitochondrial genome.
12. Regulacija genske ekspresije.	12. Regulation of gene expression.
13. Epigenetika, DNA metilacija.	13. Epigenetics, DNA methylation.
14. Rekombinantna DNA tehnologija, genetski inžiniring, biotehnologija.	14. Recombinant DNA technology, genetic engineering, biotechnology.
15. Genetska ranznolikost med posamezniki in populacijami, polimorfizmi, genetsko neravnostežje.	15. Genetic diversity between individuals and in populations, polymorphisms, linkage disequilibrium.
16. Kompleksne bolezni, multifaktorsko dedvanje.	16. Complex diseases, multifactorial inheritance.
17. Citogenetika in molekularna genetika raka.	17. Cytogenetics and molecular genetic of cancer.
18. Genetika imunskega sistema.	18. Genetics of immune system.
19. Razvojna genetika.	19. Development genetics.
20. Izbrani primeri genetskih bolezni.	20. Genetic diseases – selected cases.
21. Genomika, proteomika, farmakogenomika, personalizirana medicina.	21. Genomics, proteomics, farmacogenomics, personalized medicine.
22. Genetsko svetovanje, prenatalna diagnostika.	22. Genetic counseling, prenatal diagnosis.
23. Genetika in družba.	23. Genetic and society.

Temeljni literatura in viri / Reading materials:

1. Ronald Cohn, Stephen Scherer, Ada Hamosh. Thompson ¸& Thompson Genetics and Genomics in Medicine, 9th ed. Elsevier, 2024, ISBN: 978-0-323-54762-8.

2. Robert L. Nussbaum, Roderick R. Mclnnes, Huntington F. Willard. Thompson & Thompson Genetics in Medicine, W.B.Saunders Company, 6th ed. 2004, ISBN 0-7216-0244-4, 7th ed. 2007, ISBN: 9781416030805.

3. Robert L. Nussbaum, Roderick R. Mclnnes, Huntington F. Willard. Thompson and Thompson Genetics in Medicine with Access 8th , Elsevier, 2016, ISBN: 9781437706963.

4. Stangler Herodež Š., Erjavec Škerget A., Zagorac A., Kokalj-Vokač N.: Navodila za vaje iz molekularne biologije in genetike: skripta. Maribor: Medicinska fakulteta, 2012.
Vsakoletne dopolnitve skripte so na voljo v knjižnici MF.

Dopolnilna literatura:

1. Andrew Read and Dian Donnai. New Clinical Genetics, 4th ed., 2020, ISBN: 978-1911510703.
2. Tom Strachan, Judith Goodship & Patrick Chinnery. Genetics and genomics in medicine, CRC, 2014, ISBN: 978-0-8153-4480-3.
3. Tom Strachan, Anneke Lucassen. Genetics and genomics in medicine, 2nd ed. CRC, 2023, ISBN: 978-0-367-49082-9.

Cilji in kompetence:

Cilj predmeta je, da študentu nudi znanje o zgradbi, organizaciji in funkciji informacijskih makromulekul, nosilcih dednega materiala ter o zakonitostih prenosa in ohranjanja genetske informacije s poudarkom na humanem genomu.

Objectives and competences:

The goal of the subject is to offer the student knowledge of the structure and function of informative macromoleculs. Of principal interest are the macromolecule complexes of DNA, RNA and protein and the processes of replication, transcription and translation and technologies involved in manipulating these

Premet ponuja študentu pregledna in nekatera poglobljena znanja o zgradbi, organizaciji ter delovanju eukariontskega genoma s poudarkom na humanem genomu. Študenta seznani z možnimi načini analize genoma in ugotavljanja genetskih napak.Študenta seznani z vlogo genetskih faktorjev pri vzrokih humanih bolezni ter prispevku pri multifaktorjalnih boleznih, s kompleksno analizo delovanja in prenosa genetske informacije ter dedovanjem. Povdarek predavanj je na aplikativni vlogi genetike v sodobnih medicinskih tehnikah, diagnostiki in genski terapiji.	molecules. The courses offer to students overvew and some selected topis about organization, function and structure of eukaryotic genom with special attantion to human genom. The student gets the knowladge of major techniques used for genom analysis, and mutation analysis. The student is introduced to the role of genetic factors in etiology of human heriditary disorders, the contribution to multifactorial diseases and understanding the comlex information of functioning and transmission of genetic information. Special attantion is given to diagnostic techniques, recombinant DNA technology and gene therapy.

Predvideni študijski rezultati:	**Intended learning outcomes:**
Znanje in razumevanje: 1. Zna narisati družinsko drevo. 2. Zna ločiti strukturno od numerčne kromosomske spremembe. 3. Zna interpretirati rezultat molekularne kariotipizacije. 4. Zna izolirati DNA molekulo iz periferne krvi in določiti njeno koncentracijo. 5. Zna izvesti verižno reakcijo s polimerazo ter določiti produkt z gelsko elektroforezo. 6. Zna določiti gensko mutacijo z metodo alelno specifične verižne reakcije s polimerazo. 7. Zna določiti frekvenco alelov v populaciji. 8. Zna določiti fazo vezave alelov, izračunati frekvenco rekombinacije in verjetnost vezave dveh lokusov (Izračunati vrednost LOD.). 9. Zna izračunati tveganje za kompleksne bolezni. Prenesljive/ključne spretnosti in drugi atributi: -	Knowledge and understanding: 1. Knowledge of drawing a family pedigree. 2. Knowledge of differences between structural and numerical chromosome aberration. 3. Knowledge of interpreting the results of molecular kariotyping. 4. Knowledge of isolation of DNA molecules from periferal blood and messuring its concentration. 5. Knowledge of polymerase chain reaction and determination of the product on gel electroforesis. 6. Student knows how to determine the genetic mutation by the method of allele-specific polymerase chain reaction 7. Student knows how to determine the frequency of alleles in a population. 8. Student knows how to determine the phase of binding alleles, calculate the frequency of recombination and the probability of linkage disequlibrium (Calculate the value of the LOD.). 9. Knows how to calculate the risk of complex diseases. Transferable/Key Skills and other attributes: -

Metode poučevanja in učenja:	**Learning and teaching methods:**
Predavanja : 50% Laboratorijske vaje: 25% Seminarji: 25%	Lectures : 50% Laboratory work : 25% Seminary work: 25%

Načini ocenjevanja:	Delež (v %) / Share (in %)	**Assessment methods:**
Način (pisni izpit, ustno izpraševanje, naloge, projekt): Seminarska naloga Laboratorijsko delo	 **10** **20**	Type (examination, oral, coursework, project): Seminar paper Laboratory work

Pisni izpit Opombe: Na izpitu se zahteva znanje pridobljeno pri vseh treh oblikah poučevanja. Izpit lahko študenti opravijo s sprotnimi kolokviji, kar je zelo priporočljivo, saj tekom predavanj utrdijo znanje, ki je potrebno za razumevanje naslednjih predavanj ter pripravo seminarjev. Sprotni kolokviji so v treh sklopih. Za priznavanje pisnega izpita, morajo biti vsi kolokviji pozitivni. Kolokviji se pišejo samo na enem roku. Študenti, ki opravijo pozitivno vse tri kolokvije, seminarske naloge in kolokvij iz vaj, se prijavijo na prvi izpitni rok v juniju, kjer se jim prizna opravljen izpit. Študenti, ki niso opravljali sprotnih kolokvijev ali katerega od njih niso opravili pozitivno, pristopijo h končnemu preverjanju znanja po opravljenem kolokviju iz vaj in oddanih seminarskih nalogah, ki je v obliki pisnega izpita. Pri pisnem izpitu je možno doseči 70 točk. Vsaj 35 točk pri pisnem delu izpita je potrebnih, da se upoštevajo še točke dosežene pri vajah in seminarjih. Minimalno skupno število doseženih točk za opravljen izpit iz predmeta je 60 od 100 možnih točk. VAJE •Na vajo se je potrebno predhodno pripraviti, Z vsemi vajami študent lahko pridobi 10 točk. •Pri vsaki vaji je potrebno oddati poročilo. Vsaka uspešno (Ne samo opravljena!) opravljena vaja se točkuje z dvema točkama. Eno točko se pridobi za uspešno izvedbo vaje, drugo pa za izpolnjeno poročilo. •Ob koncu vaj je OBVEZNI kolokvij, ki se točkuje z maksimalno 10 točkami. •Skupno torej lahko študent pridobi na vajah 20 točk. •Uspešno opravljene vaje zahtevajo minimalno 12,5 točk. •Kolokvij se opravlja takoj po končanih vajah in izjemoma pred jesenskim izpitnim obdobjem (konec avgusta). SEMINARSKA NALOGA Oceno seminarja sestavlja število pravilnih odgovorov pri nalogah, ocena poročila in/oz.	**70**	Written exam Notes: On examination the knowledge acquired in all three forms of teaching is required. Students can pass the exam with colloquia which is highly recommended, because during the lectures consolidate knowledge is necessary for understanding the following lectures and seminar preparation. Colloquia are divided in three sets. For the recognition of written examination must all be positive. Colloquia are written only in one period of time! Students who pass the three colloquia, laboratory and seminary work can apply for the first examination period in June, where their exam is recognized as final examination. Students who are not engaged in ongoing colloquia or any of them have not been performed positively, accede to the final examination after completion of the laboratory and seminary work. At final exam is possible to achieve 70 points. At least 35 points in the written part of the exam is necessary to take into account the points achieved in tutorials and seminars. Minimum number of points scored by examinations of the course is 60 out of 100 possible points. LABORATORY WORK With all laboratory work the student can obtain 10 points. Each course is required to submit a report. Each successfully completed course is scored by 2 points. One point is obtained for the successful implementation of the exercise, the other for the completed report. At the end of the courses is mandatory colloquium, which counts with a maximum of 10 points. The total, therefore, a student can obtain in laboratory work is 20 points. Completion of assignments require a minimum of 12.5 points. The colloquium is carried out immediately after completion of courses and exceptionally before the autumn exam period (end of August). SEMINAR PAPER Assessment of the seminar consists of the number of correct answers in the tasks, evaluation reports and / or. rating participate in the discussion.

ocena sodelovanja v diskusiji. Seminarska naloga pomeni 10 % končne ocene pri predmetu.		Seminar paper represents 10 % of final grade for the course.

Reference nosilca / Course coordinator's references:

ANDREJAŠIČ, Nastja, BLEJC NOVAK, Anja, MOČNIK, Mirjam, MARČUN-VARDA, Nataša, STANGLER HERODEŽ, Špela, KRGOVIĆ, Danijela, ZUPAN, Andrej, MEGLIČ, Anamarija. A 15-year experience highlighting the spectrum of Alport kidney disease in the pediatric population and novel genetic variants in COL4A3-5. *Pediatric nephrology*. 2025, [v tisku][9 str.], ilustr. ISSN 1432-198X. https://link.springer.com/article/10.1007/s00467-025-06683-8, https://doi.org/10.1007/s00467-025-06683-8, DOI: 10.1007/s00467-025-06683-8. [COBISS.SI-ID 225255171], [JCR, SNIP, WoS, Scopus]
projekt: P3-0054 Patologija in molekularna genetika; financer: ARIS

VOKAČ, Damijan, STANGLER HERODEŽ, Špela, KRGOVIĆ, Danijela, KOKALJ-VOKAČ, Nadja. The role of next-generation sequencing in the management of patients with suspected non-ischemic cardiomyopathy after syncope or termination of sudden arrhythmic death. *Genes*. Jan. 2024, vol. 15, issue 1, str. [1]-11, ilustr. ISSN 2073-4425. https://doi.org/10.3390/genes15010072, https://www.mdpi.com/2073-4425/15/1/72, DOI: 10.3390/genes15010072. [COBISS.SI-ID 181229059], [JCR, SNIP, WoS, Scopus]
projekt: Financer: University Medical Centre Maribor, grant number IRP-2015/01-07
projekt: P4-220 Primerjalna genomika in genomska biodiverziteta [P4-0220]; financer: ARIS

RIHAR, Nika, KRGOVIĆ, Danijela, KOKALJ-VOKAČ, Nadja, STANGLER HERODEŽ, Špela, ZORC, Minja, DOVČ, Peter. Identification of potentially pathogenic variants for autism spectrum disorders using gene-burden analysis. *PloS one*. 2023, vol. 18, iss. 5, [article no.] e0273957, str. 1-17, ilustr. ISSN 1932-6203. https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0273957, Repozitorij Univerze v Ljubljani – RUL, DOI: 10.1371/journal.pone.0273957. [COBISS.SI-ID 152010499], [JCR, SNIP, WoS, Scopus do 1. 3. 2024: št. citatov (TC): 1, čistih citatov (CI): 1, čistih citatov na avtorja (CIAu): 0.17]

KRGOVIĆ, Danijela, GORENJAK, Mario, RIHAR, Nika, OPALIČ, Iva, STANGLER HERODEŽ, Špela, GREGORIČ KUMPERŠČAK, Hojka, DOVČ, Peter, KOKALJ-VOKAČ, Nadja. Impaired neurodevelopmental genes in Slovenian autistic children elucidate the comorbidity of autism with other developmental disorders. *Frontiers in molecular neuroscience*. Jun. 2022, vol. 15, str. 1-17, ilustr. ISSN 1662-5099. https://doi.org/10.3389/fnmol.2022.912671, https://www.frontiersin.org/articles/10.3389/fnmol.2022.912671/full, DOI: 10.3389/fnmol.2022.912671. [COBISS.SI-ID 112881155], [JCR, SNIP, WoS do 17. 7. 2023: št. citatov (TC): 1, čistih citatov (CI): 1, čistih citatov na avtorja (CIAu): 0.13, Scopus do 22. 7. 2023: št. citatov (TC): 1, čistih citatov (CI): 1, čistih citatov na avtorja (CIAu): 0.13]
projekt: Z3-9294 Advanced genomic analyses of Slovenian children with autistic spectrum disorders; financer: ARRS
projekt: P4-0220 Comparative genomics and genome biodiversity; financer: ARRS

STANGLER HERODEŽ, Špela, KOKALJ-VOKAČ, Nadja, ROŠKAR, Zlatko, DREISINGER, Mojca. Natančnost in hitrost poročanja o molekularnem odzivu pri KML bolnikih z in vitro diagnostičnim testom Xpert BCR-ABL = Accuracy and speed of molecular response reporting with Xpert BCR-ABL Ultra in vitro diagnostic test in CML patients. *Acta medico-biotechnica : AMB*. [Tiskana izd.]. 2021, vol. 14, [no.] 1, str. 49-55. ISSN 1855-5640. https://journals.um.si/index.php/amb/article/view/1428, https://dk.um.si/IzpisGradiva.php?id=83671, Digitalna knjižnica Slovenije - dLib.si, Digitalna knjižnica Univerze v Mariboru – DKUM, DOI: 10.18690/actabiomed.215. [COBISS.SI-ID 68177923]

STANGLER HERODEŽ, Špela, MARČUN-VARDA, Nataša, KOKALJ-VOKAČ, Nadja, KRGOVIĆ, Danijela. De novo KMT2D heterozygous frameshift deletion in a newborn with a congenital heart anomaly. *Balkan journal of medical genetics*. 2020, vol. 23, issue 1, str. 83-90, ilustr. ISSN 2199-5761. https://content.sciendo.com/view/journals/bjmg/23/1/article-p83.xml, https://doi.org/10.2478/bjmg-2020-0008, DOI: 10.2478/bjmg-2020-0008. [COBISS.SI-ID 28473347], [JCR, SNIP, WoS do 23. 1. 2024: št. citatov (TC): 8, čistih citatov (CI): 8, čistih citatov na avtorja (CIAu): 2.00, Scopus do 3. 1. 2024: št. citatov (TC): 7, čistih citatov (CI): 7, čistih citatov na avtorja (CIAu): 1.75]

JANČAR, Maja, KRGOVIĆ, Danijela, GORENJAK, Mario, STANGLER HERODEŽ, Špela, KOKALJ-VOKAČ, Nadja. The use of bioinformatic programming tool to analyse the data obtained with next-generation sequencing = Uporaba bioinformatskih orodij za analizo podatkov, dobljenih z metodo sekvenciranja nove generacije. V: DOVČ, Peter (ur.), ZORC, Minja (ur.). *Book of proceedings : 8th Colloquium on Genetics : 28th September 2020, online event*. [Ljubljana]: Genetic Society Slovenia, 2020. Str. 24-29. https://sgd.si/docs/PROCEEDINGS_SGD_2020.pdf. [COBISS.SI-ID 33197827]

STANGLER HERODEŽ, Špela. Določanje nukleotidnega zaporedja odsekov DNA v ginekološki onkologiji z naslednjo generacijo sekvenciranja = Determination of DNA nucleotide variations in gynecological oncology

with next-generation sequencing. V: TAKAČ, Iztok (ur.), LUKANOVIČ, Adolf (ur.), ANZELJC, Veronika. *Pregled sodobne ginekologije in porodništva pred epidemijo covida-19 leta 2020 = A state-of-the-art 2020 review on obstetrics and gynecology in the pre-Covid-19 era*. 1. izd. Maribor: Univerza v Mariboru, Univerzitetna založba: = University of Maribor, University Press, 2021. Str. 663-669, ilustr. ISBN 978-961-286-532-0. [COBISS.SI-ID 102192899]

STANGLER HERODEŽ, Špela. Molekularnogenetske preiskave v ginekološki onkologiji. V: TAKAČ, Iztok (ur.), ARKO, Darja. *Ginekološka onkologija*. 1. izd. Maribor: Univerzitetna založba Univerze, 2020. Str. 271-278. ISBN 978-961-286-330-2. [COBISS.SI-ID 512976184]