

EUROPEAN FORM
OF
AUTOBIOGRAPHY



PERSONAL INFORMATION

Name **DARINA LYUDMILOVA KACHAKOVA-YORDANOVA**
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Nationality Bulgarian
Date of birth 17.12.1984

WORK INTERNSHIP

- Dates (from-to) **March 2013 till now**
- Name and address of employer Laboratory of Genomic diagnostics, Molecular medicine center, Department of Medical chemistry and biochemistry, Medical faculty, Medical university - Sofia
- Type of activity or field of work Genetic analysis
- Position **Biology-geneticist**
- Main activities and responsibilities Planning and conducting scientific research and experiments and analyses, interpretation of obtained genetic results. Improvement or development of laboratory methods. Applying to contests for financing research projects. Conducting analyzes to determine the genetic cause of the development of a number of diseases.

- Dates (from-to) **March 2010 to March 2013**
- Name and address of employer Laboratory of Genomic diagnostics, Molecular medicine center, Department of Medical chemistry and biochemistry, Medical faculty, Medical university - Sofia
- Type of activity or field of work Genetic analysis
- Position **PhD student** registered with Order №P.185/15.03.2010
- Main activities and responsibilities Performing genetic, epigenetic, expression analyzes in prostate cancer.

- Dates (from-to) **June 2009- to March 2010**
- Name and address of employer Laboratory of Genomic diagnostics, Molecular medicine center, Department of Medical chemistry and biochemistry, Medical faculty, Medical university - Sofia
- Type of activity or field of work
- Position **Biology- laboratory assistant**

- Main activities and responsibilities Isolation of high molecular weight DNA and RNA. Participation in genetic analyses
 - Dates (from-to) **March 2009- to may 2009**
 - Name and address of employer National Genetic Laboratory, Sofia
 - Type of activity or field of work
 - Position **Medical –laboratory assistant**
- Main activities and responsibilities Isolation of high molecular weight DNA from venous blood, amniotic fluid, chorionic villi, abortive material..

EDUCATION AND TRAINING

- Dates (from-to) 2003 year- 2007year]
- Name and type of training or educational organization Sofia University „Sv. Kliment Ohridski”
 - Core subjects/preferred professional skills Специалност Молекулярна биология
- Name of the acquired qualification Bachelor

- Dates (from-to) 2007 2009 year.
- Name and type of training or educational organization Sofia University „Sv. Kliment Ohridski ”
 - Core subjects/preferred professional skills Specialty Genetics
- Name of the acquired qualification Master's degree

- Dates (from-to) 15.03.2010 to 2013 year.
- Name and type of training or educational organization Medical University - Sofia, Department of Medical Chemistry and Biochemistry
 - Core subjects/preferred professional skills Scientific specialty Molecular genetics
- Name of the acquired qualification PhD, acquired on 01.06.2015, thesis "Molecular profiling in prostate cancer"

Trainings:

2008r.- Course in Bioethics, Perugia, Italy

Camara D., Dimitrova Ir., Doynova M., Jachacz L., Kachakova D., Kepka M., Ould Isselmou CB., Vorniere JP., Yungarva Tsv. Transgenic and cloned animals: Ethical Problems? pdfs.semanticscholar.org

11.09.2008 - PALM MicroBeam LMD laser microdissection system training course, Carl Zeiss, Germany, led by Carl Zeiss representative Dr. Wolf-Dieter Schulz, Germany, held at the Center for Molecular Medicine, Department of Medicinal Chemistry and Biochemistry, Medical University of Sofia

1 - 5.11. 2010- Course "Biomek Continuum - Basic Training" for operators of automated robotic system Biomek, Nyon, Switzerland, organized by the manufacturer, Beckman, Medical University - Sofia, Department of "Medical Chemistry and Biochemistry",

7 - 9.03.2011 - Specialized course for those trained in molecular pathology of cancer: 1st EACR-OECI Joint Training Course "Molecular pathology Approach to cancer", Amsterdam

4-7.07.2011 - Specialized training on working with the PALM MicroBeam LMD laser microdissection system, Carl Zeiss, Germany and its application in the analysis of tumors, cells and cytogenetic preparations, conducted by Carl Zeiss representative Dr. Kirsten Haden-Mann, Center for Molecular Medicine

25-27.10.2011 - Bioinformatics course for working with the public database ENSEMBL and public data for: Analysis of nucleotide sequences - Nucleotide Sequence databases; Transcriptomics and Gene Expression Atlas -

ArrayExpress & Gene Expression Atlas; Mass Spectrometry and Proteomics - Mass Spectrometry based Proteomics, PRIDE and programs like BioMart for database analysis and data mining.

Organizers: the Center for Molecular Medicine and the Genome Center of the University of St. Kliment Ohridski", together with the European Bioinformatics Institute EMBL (European Bioinformatics Institute). Guest speakers from EMBL were Yana Vandrovkova, James Watson, Emma Hastings and Juan Vizqueino.

28-30.04.2014- Training on microRNA microarrays and specialized GeneSpring software for microarray data analysis, hosted by Agilent Technologies, Inc, Dr. Andreas Pölten, Waldbronn, Germany

20.04. 2022 – virtual training for introduction to analysis with DRAGEN bio IT (Map and Align, Germline pipeline, Copy Number Variant pipeline, QC Metrics (Live-demo), Accessing DRAGEN using Command Line Interface, DRAGEN Support Resources

16-17 june 2022– training for the operation and safe operation of the NovaSeq 6000, working with a server bioinformatics platform for ultra-fast secondary analysis of next-generation sequencing data - DRAGEN bio IT, as well as working with the TruSeq DNA PCR free protocol to prepare libraries for whole-genome next-generation sequencing at the reference center of the manufacturer Illumina inc - Illumina Cambridge Center at 19 Granta Park, Cambridge, United Kingdom

22-24 june 2022 online training, introduction to next-generation sequencing data analysis. June 22- Sequencing workflow, assessment of sequencing progress through SAV. Network and hardware considerations for data analysis; June 23- Introduction to FASTQ files, Secondary Analysis; June 24 – QC of sequencing data via secondary analysis metrics, Tertiary analysis. Training provided by Illumina representatives

PERSONAL SKILLS AND COMPETENCES

Acquired in life or in the profession, but not necessarily certified by an official document or diploma.

NATIVE LANGUAGE

Bulgarian

OTHER LANGUAGES

- Reading
- Writing
- Speaking

ENGLISH

EXCELLENT

GOOD

GOOD

SKILLS AND COMPETENCES

Coexistence with other people in an intercultural environment, in situations where communication and teamwork are essential (e.g. in culture and sports), etc.

ISOLATION OF DNA AND RNA FROM VARIOUS BIOLOGICAL MATERIALS, ANALYSIS OF DNA/RNA ON AGAROSE GEL, PCR TECHNOLOGIES, REAL TIME PCR, RFLP, SSCP, FRAGMENT ANALYSIS, DIRECT SEQUENCE, NEXT GENERATION SEQUENCE, ANALYSIS AND INTERPRETATION OF RESULTS OF NEXT GENERATION SEQUENCE, , MICROCHIP ANALYSIS, WORKING WITH THE BIOMEK AUTOMATED ROBOTIC SYSTEM, WORKING WITH THE PALM MICROBEAM LMD LASER MICRODISSECTION SYSTEM, ASSOCIATION STUDIES AND STATISTICS, WORKING WITH DATABASES

DRIVING LICENSE

Driving license for category B and M motor vehicles

ADDITIONAL INFORMATION

REFERRALS WILL BE PROVIDED UPON REQUEST.

Individuals and/or organizations that can provide a professional recommendation or review:

1. Prof. Alexey Savov, Department of Obstetrics and Gynecology, Faculty of Medicine, MU
2. Prof. Dr. Radka Kaneva, Department of Biochemistry and Chemistry, Faculty of Medicine, MU

APPLICATIONS

LIST OF SCIENTIFIC PUBLICATIONS

COPY OF THE WORKBOOK

MEDICAL CERTIFICATE

CRIMINAL RECORD CERTIFICATE

MASTER'S DIPLOMA NOTARIZED

Application 1

List of scientific publications of Darina Kachakova

1. Eeles RA, Kote-Jarai Z, Olama AA, Kaneva R, Slavov C, Mitkova A, Kachakova D..., The UK Genetic Prostate Cancer Study Collaborators/British Association of Urological Surgeons' Section of Oncology, The UK ProtecT Study Collaborators, The PRACTICAL Consortium, Easton DF, Identification of seven novel prostate cancer susceptibility loci through a genome-wide association study, *Nature Genetics* 2009, Oct;41(10):1116-21., october 2009; IF 2009 (34.284)
2. Kote-Jarai, Sofia, Ali Amin Al Olama, Graham G. Danielle M. Karyadi ... Mitev...The **PRACTICAL** Consortium... (Kachakova D)...., Douglas F Easton, Rosalind A Eeles. The Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. *Nat Genet.* 2011 Jul 10. doi: 10.1038/ng.882. PubMed PMID: 21743467; ИФ 2011 (22.84)
3. Amin Al Olama A, Kote-Jarai Z, Schumacher FR, Wiklund F, Berndt SI, Benlloch S, Giles GG,Cybulski C, Lubinski J, Thibodeau SN, Schaid DJ, Sorensen KD, Batra J, Clements JA, Chambers S, Aitken J, Gardiner RA, Maier C, Vogel W, Dörk T, Brenner H, Habuchi T, Ingles S, John EM, Dickinson JL, Cannon-Albright L, Teixeira MR, Kaneva R, Zhang HW, Lu YJ, Park JY, Cooney KA, Muir KR, Leongamornlert DA, Saunders E, Tymrakiewicz M, Mahmud N, Guy M, Govindasami K, O'Brien LT, Lose F, McDonnell SK, Joshi AD, Shahabi A, Pinto P, Santos J, Ray A, Sellers TA, Lin HY, Stephenson RA, Teerlink C, Muller H, Rothenbacher D, Tsuchiya N, Narita S, Cao GW, Slavov C, Mitev V; The UK Genetic Prostate Cancer Study Collaborators/British Association of Urological Surgeons' Section of Oncology; The UK ProtecT Study Collaborators; The Australian Prostate Cancer Bioresource; The **PRACTICAL** Consortium (... , Kachakova D, Mitkova A, Goranova T, Stancheva G,....), Chanock S, Gronberg H, Haiman CA, Kraft P, Easton DF, Eeles RA. A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. *Hum Mol Genet.* 2013 Jan 15;22(2):408-15; PMID: IF 2013 (6.677)
4. Giles GG, Severi G, Wiklund F, Gronberg H, Haiman CA, Schumacher F, Henderson BE, Le Marchand L, Lindstrom S, Kraft P, Hunter DJ, Gapstur S, Chanock S, Berndt SI, Albanes D, Andriole G, Schleutker J, Weischer M, Canzian F, Riboli E, Key TJ, Travis RC, Campa D, Ingles SA, John EM, Hayes RB, Pharoah P, Khaw KT, Stanford JL, Ostrander EA, Signorello LB, Thibodeau SN, Schaid D, Maier C, Vogel W, Kibel AS, Cybulski C, Lubinski J, Cannon-Albright L, Brenner H, Park JY, Kaneva R, Batra J, Spurdle A, Clements JA, Teixeira MR, Govindasami K, Guy M, Wilkinson RA, Sawyer EJ, Morgan A, Dicks E, Baynes C, Conroy D, Bojesen SE, Kaaks R, Vincent D, Bacot F, Tessier DC; COGS-CRUK GWAS-ELLIPSE (Part of GAME-ON) Initiative; UK Genetic Prostate Cancer Study Collaborators/British Association of Urological Surgeons' Section of Oncology; UK ProtecT Study Collaborators; **PRACTICAL** Consortium, Easton DF, Eeles RA. (2013) Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. *Hum Mol Genet.* 2013 Jun 15;22(12):2520-8. IF 2013 (6.677)
5. **Kachakova D**, Mitkova A, Popov E, Beltcheva O, Vlahova A, Dikov T, Hristova S, Mitev V, Slavov C, **Kaneva R**. Evaluation of the clinical value of the newly identified urine biomarker HIST1H4K for diagnosis and prognosis of prostate cancer in Bulgarian patients. *J BUON.* 2013 Jul-Sep;18(3):660-8. IF 2013 (0.706)
6. Popov TM, Stancheva I, **Kachakova DL**, Rangachev J, Konov D, Varbanova S, Mitev VI, **Kaneva RP**, Popova DP. Auditory Outcome After Cochlear Implantation in Patients With

7. **Д. Качакова**, А. Миткова, Радка Кънева, Ваньо Митев. Ракът на простатата-генетично, геномно и епигенетично заболяване. Биоамаркери. *Studia Oncologica*, Октомври 2014, година VI, брой 3

8. Marinova, D., Slavova, Y., **Kachakova, D.**, Stancheva, G., Mitkova, A., Kaneva, R., ... & Mitev, V. (2014). Gene expression of EGFR, MINA53, MEN1 and MTOR in NSCLCs. *European Respiratory Journal*, 44(Suppl 58), P2709.

9. **Kachakova D**, Mitkova A, Popov E, Popov I, Vlahova A, Dikov T, Christova S, Mitev V, Slavov C, Kaneva R. Combinations of Serum Prostate-Specific Antigen and Plasma Expression Levels of let-7c, miR-30c, miR-141, and miR-375 as Potential Better Diagnostic Biomarkers for Prostate Cancer. *DNA Cell Biol.* 2015 Mar;34(3):189-200, IF 2015 (2.574)

10. Kote-Jarai Z, Saunders EJ, Leongamornlert DA, Tymrakiewicz M, Dadaev T, Jugurnauth-Little S, Ross-Adams H, Al Olama AA, Benlloch S, Halim S, Russell R, Dunning AM, Luccarini C, Dennis J, Neal DE, Hamdy FC, Donovan JL, Muir K, Giles GG, Severi G, Wiklund F, Gronberg H, Haiman CA, Schumacher F, Henderson BE, Le Marchand L, Lindstrom S, Kraft P, Hunter DJ, Gapstur S, Chanock S, Berndt SI, Albanes D, Andriole G, Schleutker J, Weischer M, Canzian F, Riboli E, Key TJ, Travis RC, Campa D, Ingles SA, John EM, Hayes RB, Pharoah P, Khaw KT, Stanford JL, Ostrander EA, Signorello LB, Thibodeau SN, Schaid D, Maier C, Vogel W, Kibel AS, Cybulski C, Lubinski J, Cannon-Albright L, Brenner H, Park JY, Kaneva R, Batra J, Spurdle A, Clements JA, Teixeira MR, Govindasami K, Guy M, Wilkinson RA, Sawyer EJ, Morgan A, Dicks E, Baynes C, Conroy D, Bojesen SE, Kaaks R, Vincent D, Bacot F, Tessier DC; COGS-CRUK GWAS-ELLIPSE (Part of GAME-ON) Initiative; UK Genetic Prostate Cancer Study Collaborators/British Association of Urological Surgeons' Section of Oncology; UK ProtecT Study Collaborators; **PRACTICAL** Consortium, Easton DF, Eeles RA. (2013) Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. *Hum Mol Genet.* 2013 Jun 15;22(12):2520-8. IF 2013 (6.677)

11. Eeles RA, Olama AA, Benlloch S, Saunders EJ, Leongamornlert DA, Tymrakiewicz M, Ghousaini M, Luccarini C, Dennis J, Jugurnauth-Little S, Dadaev T, Neal DE, Hamdy FC, Donovan JL, Muir K, Giles GG, Severi G, Wiklund F, Gronberg H, Haiman CA, Schumacher F, Henderson BE, Le Marchand L, Lindstrom S, Kraft P, Hunter DJ, Gapstur S, Chanock SJ, Berndt SI, Albanes D, Andriole G, Schleutker J, Weischer M, Canzian F, Riboli E, Key TJ, Travis RC, Campa D, Ingles SA, John EM, Hayes RB, Pharoah PD, Pashayan N, Khaw KT, Stanford JL, Ostrander EA, Signorello LB, Thibodeau SN, Schaid D, Maier C, Vogel W, Kibel AS, Cybulski C, Lubinski J, Cannon-Albright L, Brenner H, Park JY, Kaneva R, Batra J, Spurdle AB, Clements JA, Teixeira MR, Dicks E, Lee A, Dunning AM, Baynes C, Conroy D, Maranian MJ, Ahmed S, Govindasami K, Guy M, Wilkinson RA, Sawyer EJ, Morgan A, Dearnaley DP, Horwich A, Huddart RA, Khoo VS, Parker CC, Van As NJ, Woodhouse CJ, Thompson A, Dudderidge T, Ogden C, Cooper CS, Lophatananon A, Cox A, Southey MC, Hopper JL, English DR, Aly M, Adolfsson J, Xu J, Zheng SL, Yeager M, Kaaks R, Diver WR, Gaudet MM, Stern MC, Corral R, Joshi AD, Shahabi A, Wahlfors T, Tammela TL, Auvinen A, Virtamo J, Klarskov P, Nordestgaard BG, Røder MA, Nielsen SF, Bojesen SE, Siddiq A, Fitzgerald LM, Kolb S, Kwon EM, Karyadi DM, Blot WJ, Zheng W, Cai Q, McDonnell SK, Rinckleb AE, Drake B, Colditz G, Wokolorczyk D, Stephenson RA, Teerlink C, Muller H, Rothenbacher D, Sellers TA, Lin HY, Slavov C, Mitev V, Lose F, Srinivasan S, Maia S, Paulo P, Lange E, Cooney KA, Antoniou AC, Vincent D, Bacot F, Tessier DC; COGS-Cancer Research UK GWAS-ELLIPSE (part of GAME-ON) Initiative; Australian Prostate

Cancer Bioresource; UK Genetic Prostate Cancer Study Collaborators/British Association of Urological Surgeons' Section of Oncology; UK ProtecT (Prostate testing for cancer and Treatment) Study Collaborators; **PRACTICAL** (Prostate Cancer Association Group to Investigate Cancer-Associated Alterations in the Genome) Consortium, Kote-Jarai Z, Easton DF. Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. *Nat Genet.* 2013 Apr;45(4):385-91, 391e1-2. doi:10.1038/ng.2560. PubMed PMID: 23535732; PubMed Central PMCID: PMC3832790. IF 2013 (29.648)

12. Al Olama AA, Kote-Jarai Z, Berndt SI, Conti DV, Schumacher F, Han Y, Benlloch S, Hazelett DJ, Wang Z, Saunders E, Leongamornlert D, Lindstrom S, Jugurnauth-Little S, Dadaev T, Tymrakiewicz M, Stram DO, Rand K, Wan P, Stram A, Sheng X, Pooler LC, Park K, Xia L, Tyrer J, Kolonel LN, Le Marchand L, Hoover RN, Machiela MJ, Yeager M, Burdette L, Chung CC, Hutchinson A, Yu K, Goh C, Ahmed M, Govindasami K, Guy M, Tammela TL, Auvinen A, Wahlfors T, Schleutker J, Visakorpi T, Leinonen KA, Xu J, Aly M, Donovan J, Travis RC, Key TJ, Siddiq A, Canzian F, Khaw KT, Takahashi A, Kubo M, Pharoah P, Pashayan N, Weischer M, Nordestgaard BG, Nielsen SF, Klarskov P, Røder MA, Iversen P, Thibodeau SN, McDonnell SK, Schaid DJ, Stanford JL, Kolb S, Holt S, Knudsen B, Coll AH, Gapstur SM, Diver WR, Stevens VL, Maier C, Luedeke M, Herkommer K, Rinckleb AE, Strom SS, Pettaway C, Yeboah ED, Tettey Y, Biritwum RB, Adjei AA, Tay E, Truelove A, Niwa S, Chokkalingam AP, Cannon-Albright L, Cybulski C, Wokołarczyk D, Kluźniak W, Park J, Sellers T, Lin HY, Isaacs WB, Partin AW, Brenner H, Dieffenbach AK, Stegmaier C, Chen C, Giovannucci EL, Ma J, Stampfer M, Penney KL, Mucci L, John EM, Ingles SA, Kittles RA, Murphy AB, Pandha H, Michael A, Kierzek AM, Blot W, Signorello LB, Zheng W, Albanes D, Virtamo J, Weinstein S, Nemesure B, Carpten J, Leske C, Wu SY, Hennis A, Kibel AS, Rybicki BA, Neslund-Dudas C, Hsing AW, Chu L, Goodman PJ, Klein EA, Zheng SL, Batra J, Clements J, Spurdle A, Teixeira MR, Paulo P, Maia S, Slavov C, Kaneva R, Mitev V, Witte JS, Casey G, Gillanders EM, Seminara D, Riboli E, Hamdy FC, Coetzee GA, Li Q, Freedman ML, Hunter DJ, Muir K, Gronberg H, Neal DE, Southey M, Giles GG, Severi G; Breast and Prostate Cancer Cohort Consortium (BPC3); **PRACTICAL** (Prostate Cancer Association Group to Investigate Cancer-Associated Alterations in the Genome) Consortium; COGS (Collaborative Oncological Gene-environment Study) Consortium; GAME-ON/ELLIPSE Consortium, Cook MB, Nakagawa H, Wiklund F, Kraft P, Chanock SJ, Henderson BE, Easton DF, Eeles RA, Haiman CA. A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. *Nat Genet.* 2014 Oct;46(10):1103-9. doi: 10.1038/ng.3094. Epub 2014 Sep 14, IF 2014 (29.648)

13. Yu OH, Foulkes WD, Dastani Z, Martin RM, Eeles R; **PRACTICAL** Consortium; CRUK GWAS Investigators, Richards JB. An assessment of the shared allelic architecture between type II diabetes and prostate cancer. *Cancer Epidemiol Biomarkers Prev.* 2013 Aug;22(8):1473-5. doi: 10.1158/1055-9965.EPI-13-0476. Epub 2013 May 23. PubMed PMID: 23704474 IF 2013 (4.324)

14. Shen H, Fridley BL, Song H, Lawrenson K, Cunningham JM, Ramus SJ, Cicek MS, Tyrer J, Stram D, Larson MC, Köbel M; **PRACTICAL** Consortium, Ziogas A, Zheng W, Yang HP, Wu AH, Wozniak EL, Woo YL, Winterhoff B, Wik E, Whittemore AS, Wentzensen N, Weber RP, Vitonis AF, Vincent D, Vierkant RA, Vergote I, Van Den Berg D, Van Altena AM, Tworoger SS, Thompson PJ, Tessier DC, Terry KL, Teo SH, Templeman C, Stram DO, Southey MC, Sieh W, Siddiqui N, Shvetsov YB, Shu XO, Shridhar V, Wang-Gohrke S, Severi G, Schwaab I, Salvesen HB, Rzepecka IK, Runnebaum IB, Rossing MA, Rodriguez-Rodriguez L, Risch HA, Renner SP, Poole EM, Pike MC, Phelan CM, Pelttari LM, Pejovic T, Paul J, Orlov I, Omar SZ, Olson SH, Odunsi K, Nickels S, Nevanlinna H, Ness RB, Narod SA, Nakanishi T, Moysich KB, Monteiro AN, Moes-Sosnowska J, Modugno F, Menon U, McLaughlin JR, McGuire V, Matsuo K, Adenan NA, Massuger LF, Lurie G, Lundvall L, Lubiński J, Lissowska J, Levine DA, Leminen A, Lee AW, Le ND, Lambrechts S, Lambrechts

D, Kupryjanczyk J, Krakstad C, Konecny GE, Kjaer SK, Kiemeny LA, Kelemen LE, Keeney GL, Karlan BY, Karevan R, Kalli KR, Kajiyama H, Ji BT, Jensen A, Jakubowska A, Iversen E, Hosono S, Høgdall CK, Høgdall E, Hoatlin M, Hillemanns P, Heitz F, Hein R, Harter P, Halle MK, Hall P, Gronwald J, Gore M, Goodman MT, Giles GG, Gentry-Maharaj A, Garcia-Closas M, Flanagan JM, Fasching PA, Ekici AB, Edwards R, Eccles D, Easton DF, Dürst M, du Bois A, Dörk T, Doherty JA, Despierre E, Dansonka-Mieszkowska A, Cybulski C, Cramer DW, Cook LS, Chen X, Charbonneau B, Chang-Claude J, Campbell I, Butzow R, Bunker CH, Brüeggmann D, Brown R, Brooks-Wilson A, Brinton LA, Bogdanova N, Block MS, Benjamin E, Beesley J, Beckmann MW, Bandera EV, Baglietto L, Bacot F, Armasu SM, Antonenkova N, Anton-Culver H, Aben KK, Liang D, Wu X, Lu K, Hildebrandt MA; Australian Ovarian Cancer Study Group; Australian Cancer Study, Schildkraut JM, Sellers TA, Huntsman D, Berchuck A, Chenevix-Trench G, Gayther SA, Pharoah PD, Laird PW, Goode EL, Pearce CL. Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. *Nat Commun.* 2013;4:1628. doi: 10.1038/ncomms2629. IF 2013 (10.742)

15. Laitinen VH, Rantapero T, Fischer D, Vuorinen EM, Tammela TL; **PRACTICAL Consortium**, Wahlfors T, Schleutker J. Fine-mapping the 2q37 and 17q11.2-q22 loci for novel genes and sequence variants associated with a genetic predisposition to prostate cancer. *Int J Cancer.* 2015 May 15;136(10):2316-27. IF 2015 (5.531)

16. Panagiotou OA, Travis RC, Campa D, Berndt SI, Lindstrom S, Kraft P, Schumacher FR, Siddiq A, Papatheodorou SI, Stanford JL, Albanes D, Virtamo J, Weinstein SJ, Diver WR, Gapstur SM, Stevens VL, Boeing H, Bueno-de-Mesquita HB, Gurrea AB, Kaaks R, Khaw KT, Krogh V, Overvad K, Riboli E, Trichopoulos D, Giovannucci E, Stampfer M, Haiman C, Henderson B, Le Marchand L, Gaziano JM, Hunter DJ, Koutros S, Yeager M, Hoover RN; The **PRACTICAL Consortium**, Chanock SJ, Wacholder S, Key TJ, Tsilidis KK. A Genome-wide Pleiotropy Scan for Prostate Cancer Risk. *Eur Urol.* 2015 Apr;67(4):649-57. doi: 10.1016/j.eururo.2014.09.020. [Epub ahead of print]; IF 2015 (14.976)

17. Han Y, Signorello LB, Strom SS, Kittles RA, Rybicki BA, Stanford JL, Goodman PJ, Berndt SI, Carpten J, Casey G, Chu L, Conti DV, Rand KA, Diver WR, Hennis AJ, John EM, Kibel AS, Klein EA, Kolb S, Le Marchand L, Leske MC, Murphy AB, Neslund-Dudas C, Park JY, Pettaway C, Rebbeck TR, Gapstur SM, Zheng SL, Wu SY, Witte JS, Xu J, Isaacs W, Ingles SA, Hsing A; **PRACTICAL Consortium**; ELLIPSE GAME-ON Consortium, Easton DF, Eeles RA, Schumacher FR, Chanock S, Nemesure B, Blot WJ, Stram DO, Henderson BE, Haiman CA. Generalizability of established prostate cancer risk variants in men of African ancestry. *Int J Cancer.* 2015 Mar 1;136(5):1210-7. doi: 10.1002/ijc.29066. Epub 2014 Jul 15. IF 2015 (5.531)

18. Andreassen OA, Zuber V, Thompson WK, Schork AJ, Bettella F; **PRACTICAL Consortium**; CRUK GWAS, Djurovic S, Desikan RS, Mills IG, Dale AM. Shared common variants in prostate cancer and blood lipids. *Int J Epidemiol.* 2014 Aug;43(4):1205-14. doi: 10.1093/ije/dyu090. Epub 2014 Apr 30. IF 2014 (9.197)

Marinova, D. M., Slavova, Y., Stancheva, G., **Kachakova, D.**, Mitkova, A., & Kaneva, R. (2014, April). GENE EXPRESSION OF EPIDERMAL GROWTH FACTOR RECEPTOR, MYC-INDUCED NUCLEAR ANTIGEN, 53KDA AND MECHANISTIC TARGET OF RAPAMYCIN IN PULMONARY NEUROENDOCRINE TUMORS. In *JOURNAL OF THORACIC ONCOLOGY* (Vol. 9, No. 4, pp. S10-S10). 530 WALNUT ST, PHILADELPHIA, PA 19106-3621 USA: LIPPINCOTT WILLIAMS & WILKINS.

19. **Kachakova D**, Mitkova A, Popov E, Beltcheva O, Vlahova A, Dikov T, Christova S, Mitev V, Slavov C, Kaneva R. Polymorphisms in androgen metabolism genes AR, CYP1B1, CYP19,

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