



The University of Alabama at Birmingham

QUARTERLY GENETICS UPDATE SPRING 2022

Researchers pioneering long-read sequencing studies explain why long reads matter



With long-read sequencing, read lengths can average 10,000 to 30,000 base pairs, with a maximum length over 1 megabase (1 million base pairs) on an Oxford Nanopore device, says **Zechen Chong, Ph.D.**, assistant professor in the Department of Genetics at the Heersink School of Medicine. Chong's lab is focused on developing new algorithms for the analysis of long-read sequencing data, and he is collaborating with several UAB researchers using long-read sequencing.

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UAB Cardiogenomics Clinic transforms cardiac care through genomic medicine



"The approach to understanding the risk of developing potentially fatal heart diseases can vary, from using the latest advances in genetics to simply talking with our grandparents," said **Ali Al-Beshri, M.D.**, an assistant professor in the UAB Department of Genetics. "As technology has evolved, we have gained an improved understanding of what to do and how to manage individuals who are at risk for inheriting cardiovascular diseases. Furthermore, the latest advances in genetic medicine such as gene-based therapeutics are becoming increasingly common in the clinical pipeline."

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Distinct structural domains in MUNC long non-coding RNA regulate gene expression



In a paper in *Cell Reports*, **Anindya Dutta, M.D., Ph.D.**, and colleagues detail the first structure-function study of a long non-coding RNA, or lncRNA, called MUNC lncRNA. Their findings point out the importance of experimentally determining the structure of an lncRNA through a chemical method — rather than computer modeling — to identify structural domains that turn out to have multiple different functions.

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How a game-changing transplant could treat dying organs



Scientists have long known that defective mitochondria can cause biological chaos. “Mitochondrial dysfunction is a universal driver of disease,” says **Keshav Singh**, a mitochondria researcher at the University of Alabama Birmingham who founded the Mitochondria Research and Medicine Society in the U.S. and India. Whether tissue damage is caused by disease or even space travel, faulty mitochondria are frequently involved.

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Robustly collaborative projects funded through annual AMC21 Multi-PI Awards; Lee, Pollock, and Korf advance cutting-edge research



The Heersink School of Medicine has awarded the annual Multi-Investigator Program Award to three research investigators for the 2021-2022 cycle: Timmy Lee, M.D., **Bruce Korf, M.D., Ph.D.**, and David Pollock, Ph.D. Dr. Korf was awarded a Multi-PI Award for his study UAB Gene Therapy Neurofibromatosis Program.

[Read more](#)

Dr. Chong selected for Featured Discovery

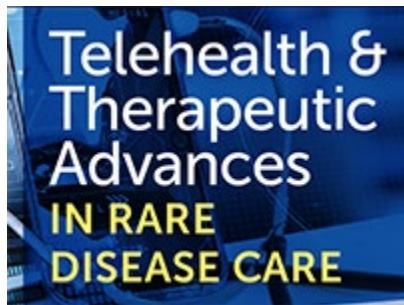


Zechen Chong, Ph.D., assistant professor in the Department of Genetics, is the latest winner of the Heersink School of Medicine's Featured Discovery. This initiative celebrates important research from Heersink faculty members.

Chong's study, "Accurate long-read de novo assembly evaluation with Inspector," was published in *Genome Biology*.

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Telehealth and therapeutic advances are focus of 2022 Rare Disease Genomics Symposium



The University of Alabama at Birmingham presented the ninth annual Rare Disease Genomics Symposium on March 4 virtually. The symposium is held every year in conjunction with World Rare Disease Day and is hosted by UAB, Children's of Alabama, Alabama Rare and the Alabama Genomic Health Initiative, and the UAB Center for Clinical and Translational Science. This year's event focused on science- and research-related telehealth and therapeutic advances. Participants heard from experts in the fields of genetics, telemedicine, precision medicine and more.

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All of Us research program releases first genomic data, creating unprecedented platform for health discoveries



The National Institutes of Health's All of Us Research Program has made its initial genomic data available, bringing nearly 100,000 whole genome sequences onto its data analysis platform, the Researcher Workbench. As the All of Us Southern Network leader with sites across Alabama,

Mississippi, and Louisiana, UAB Medicine has played a significant role in this milestone, nurturing relationships with diverse participants from throughout the region to ensure substantial diversity in the dataset.

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Recent Faculty Publications

Gooch C, Souder JP, Tedder ML, Kerkhof J, Lee JA, Louie RJ, Sadikovic B, Fletcher RS, **Robin NH**. Near complete deletion of KMT2D in a college student. *Am J Med Genet A*. 2022 Jan 18. doi: 10.1002/ajmg.a.62652. Online ahead of print.

Korf B. An interview on rare and genetic diseases with Dr Bruce Korf, Associate Dean for Genomic Medicine at the University of Alabama at Birmingham. *Curr Med Res Opin*. 2022 Feb;38(2):161-163. doi: 10.1080/03007995.2021.2023395. Epub 2022 Jan 20.

Yan D, Qin H, Wu HY, **Chen JY**. Editorial: AI-Enabled Data Science for COVID-19. *Front Big Data*. 2021 Nov 24;4:802452. doi: 10.3389/fdata.2021.802452. eCollection 2021.

Chen Y, Zhang Y, Wang AY, Gao M, **Chong Z**. Accurate long-read de novo assembly evaluation with Inspector. *Genome Biol*. 2021 Nov 14;22(1):312. doi: 10.1186/s13059-021-02527-4.

Zhao Q, Yan S, Lu J, Parker DJ, Wu H, Sun Q, **Crossman DK**, Liu S, Wang Q, Sesaki H, **Mitra K**, Liu K, **Jiao K**. Drp1 regulates transcription of ribosomal protein genes in embryonic hearts. *J Cell Sci*. 2022 Feb 15;135(4):jcs258956. doi: 10.1242/jcs.258956. Epub 2022 Feb 21.

Awad EK, Moore M, Liu H, Ciszewski L, **Lambert L**, **Korf BR**, Popplewell L, **Kesterson RA**, **Wallis D**. Restoration of Normal NF1 Function with Antisense Morpholino Treatment of Recurrent Pathogenic Patient-Specific Variant c.1466A>G; p.Y489C. *J Pers Med*. 2021 Dec 7;11(12):1320. doi: 10.3390/jpm11121320.

Davenport ML, Davis MR, Davenport BN, **Crossman DK**, Hall A, Pike J, Harada S, Hurst DR, **Edmonds MD**. Suppression of SIN3A by miR-183 promotes breast cancer metastasis. *Mol Cancer Res*. 2022 Mar 1:molcanres.0508.2021. doi: 10.1158/1541-

7786.MCR-21-0508. Online ahead of print.

Cui X, Zhang C, Xu Z, Wang S, Li X, Stringer-Reasor E, Bae S, Zeng L, Zhao D, **Liu R**, Qi LS, **Wang L**. Dual CRISPR interference and activation for targeted reactivation of X-linked endogenous FOXP3 in human breast cancer cells. *Mol Cancer*. 2022 Feb 7;21(1):38. doi: 10.1186/s12943-021-01472-x.

Wang C, Leavenworth J, Zhang C, Liu Z, Yuan KY, Wang Y, Zhang G, Wang S, Cui X, Zhang Y, Bae S, Zhou J, **Wang L**, **Liu R**. Epigenetic regulation of EIF4A1 through DNA methylation and an oncogenic role of eIF4A1 through BRD2 signaling in prostate cancer. *Oncogene*. 2022 Mar 31. doi: 10.1038/s41388-022-02272-3. Online ahead of print.

Hyndman KA, **Crossman DK**. Kidney cell type-specific changes in the chromatin and transcriptome landscapes following epithelial Hdac1 and Hdac2 knockdown. *Physiol Genomics*. 2022 Feb 1;54(2):45-57. doi: 10.1152/physiolgenomics.00102.2021. Epub 2021 Dec 10.

Küry S, Ebstein F, Mollé A, Besnard T, Lee MK, Vignard V, Hery T, Nizon M, Mancini GMS, Giltay JC, Cogné B, McWalter K, Deb W, Mor-Shaked H, Li H, Schnur RE, Wentzensen IM, Denommé-Pichon AS, Fourgeux C, Verheijen FW, Faurie E, Schot R, Stevens CA, Smits DJ, Barr E, Sheffer R, Bernstein JA, Stimach CL, Kovitch E, Shashi V, Schoch K, Smith W, van Jaarsveld RH, **Hurst ACE**, Smith K, Baugh EH, Bohm SG, Vyhnálková E, Ryba L, Delnatte C, Neira J, Bonneau D, Toutain A, Rosenfeld JA; Undiagnosed Diseases Network, Audebert-Bellanger S, Gilbert-Dussardier B, Odent S, Laumonnier F, Berger SI, Smith ACM, Bourdeaut F, Stern MH, Redon R, Krüger E, Margueron R, Bézieau S, Poschmann J, Isidor B. Rare germline heterozygous missense variants in BRCA1-associated protein 1, BAP1, cause a syndromic neurodevelopmental disorder. *Am J Hum Genet*. 2022 Feb 3;109(2):361-372. doi: 10.1016/j.ajhg.2021.12.011. Epub 2022 Jan 19.

Kontaridis MI, Roberts AE, Schill L, Schoyer L, Stronach B, Andelfinger G, Aoki Y, Axelrad ME, Bakker A, Bennett AM, Broniscer A, Castel P, Chang CA, Cyganek L, Das TK, den Hertog J, Galperin E, Garg S, Gelb BD, Gordon K, Green T, Gripp KW, Itkin M, Kiuru M, **Korf BR**, Livingstone JR, López-Juárez A, Magoulas PL, Mansour S, Milner T, Parker E, Pierpont EI, Plouffe K, Rauen KA, Shankar SP, Smith SB, Stevenson DA, Tartaglia M, Van R, Wagner ME, Ware SM, Zenker M. The seventh international RASopathies symposium: Pathways to a cure-expanding knowledge, enhancing research, and therapeutic discovery. *Am J Med Genet A*. 2022 Mar 9. doi: 10.1002/ajmg.a.62716. Online ahead of print.

Winter SS, McCaustland A, Qu C, Simeona N, Heerema NA, **Carroll AJ**, Wood BL, Gheorghe G, Mullighan CG, Wilson BS. VpreB surrogate light chain expression in B-lineage ALL: a report from the Children's Oncology Group. *Blood Adv*. 2022 Jan 25;6(2):585-589. doi: 10.1182/bloodadvances.2021005245.

Fobare S, Kohlschmidt J, Ozer HG, Mrózek K, Nicolet D, Mims AS, Garzon R, Blachly JS, Orwick S, **Carroll AJ**, Stone RM, Wang ES, Kolitz JE, Powell BL, Oakes CC, Eisfeld AK, Hertlein E, Byrd JC. Molecular, clinical, and prognostic implications of PTPN11

mutations in acute myeloid leukemia. *Blood Adv.* 2022 Mar 8;6(5):1371-1380. doi: 10.1182/bloodadvances.2021006242.

Banka S, Bennington A, Baker MJ, Rijckmans E, Clemente GD, Ansor NM, Sito H, Prasad P, Anyane-Yeboa K, Badalato L, Dimitrov B, Fitzpatrick D, **Hurst ACE**, Jansen AC, Kelly MA, Krantz I, Rieubland C, Ross M, Rudy NL, Sanz J, Stouffs K, Xu ZL, Malliri A, Kazanietz MG, Millard TH. Activating RAC1 variants in the switch II region cause a developmental syndrome and alter neuronal morphology. *Brain.* 2022 Feb 9:awac049. doi: 10.1093/brain/awac049. Online ahead of print.

Gupta S, Teachey DT, Chen Z, Rabin KR, Dunsmore KP, Larsen EC, Maloney KW, Mattano LA Jr, Winter SS, **Carroll AJ**, Heerema NA, Borowitz MJ, Wood BL, Carroll WL, Raetz EA, Winick NJ, Loh ML, Hunger SP, Devidas M. Sex-based disparities in outcome in pediatric acute lymphoblastic leukemia: a Children's Oncology Group report. *Cancer.* 2022 Feb 24. doi: 10.1002/cncr.34150. Online ahead of print.

Arend RC, Scalise CB, Gordon ER, Davis AM, Foxall ME, Johnston BE, **Crossman DK**, Cooper SJ. Metabolic Alterations and WNT Signaling Impact Immune Response in HGSC. *Clin Cancer Res.* 2022 Apr 1;28(7):1433-1445. doi: 10.1158/1078-0432.CCR-21-2984.

Dominguez AA, Chavez MG, Urke A, Gao Y, **Wang L**, Qi LS. CRISPR-mediated Synergistic Epigenetic and Transcriptional Control. *CRISPR J.* 2022 Mar 10. doi: 10.1089/crispr.2021.0099. Online ahead of print.

Kaza N, Lin VY, Stanford D, Hussain SS, Libby EF, Kim H, Borgonovi M, Conrath K, Mutyam V, Byzek SA, Tang LP, Trombley JE, Rasmussen L, **Schoeb T**, Leung HM, Tearney GJ, Raju SV, Rowe SM. Evaluation of a novel CFTR potentiator in cpd ferrets with acquired cftr dysfunction. *Eur Respir J.* 2021 Dec 16:2101581. doi: 10.1183/13993003.01581-2021. Online ahead of print.

Deng J, Hartung T, Capobianco E, **Chen JY**, Emmert-Streib F. Editorial: Artificial Intelligence for Precision Medicine. *Front Artif Intell.* 2022 Jan 21;4:834645. doi: 10.3389/frai.2021.834645. eCollection 2021.

Chen J, Li F, Wang M, Li J, **Marquez-Lago TT**, **Leier A**, Revote J, Li S, Liu Q, Song J. BigFiRSt: A Software Program Using Big Data Technique for Mining Simple Sequence Repeats From Large-Scale Sequencing Data. *Front Big Data.* 2022 Jan 18;4:727216. doi: 10.3389/fdata.2021.727216. eCollection 2021.

Vu T, Datta A, Banister C, Jin L, Yuan G, Samuel T, Bae S, Eltoum IE, Manne U, Zhang B, Welner RS, **Mitra K**, Buckhaults P, Datta PK. Serine-threonine Kinase Receptor-Associated Protein is a Critical Mediator of APC Mutation-Induced Intestinal Tumorigenesis Through a Feed-Forward Mechanism. *Gastroenterology.* 2022 Jan;162(1):193-208. doi: 10.1053/j.gastro.2021.09.010. Epub 2021 Sep 11.

Bowling KM, Thompson ML, Finnila CR, Hiatt SM, Latner DR, Amaral MD, Lawlor JM, East KM, Cochran ME, Greve V, Kelley WV, Gray DE, Felker SA, Meddaugh H, Cannon A, Luedcke A, Jackson KE, Hendon LG, Janani HM, Johnston M, Merin LA, Deans SL,

Tuura C, Williams H, Laborde K, Neu MB, Patrick-Esteve J, **Hurst ACE**, Kandasamy J, Carlo W, Brothers KB, Kirmse BM, Savich R, Superneau D, Spedale SB, Knight SJ, Barsh GS, **Korf BR**, Cooper GM. Genome sequencing as a first-line diagnostic test for hospitalized infants. *Genet Med.* 2021 Nov 27:S1098-3600(21)05400-9. doi: 10.1016/j.gim.2021.11.020. Online ahead of print.

Isidor B, Ebstein F, **Hurst A**, Vincent M, Bader I, Rudy NL, Cogne B, Mayr J, Brehm A, Bupp C, Warren K, Bacino CA, Gerard A, Ranells JD, Metcalfe KA, van Bever Y, Jiang YH, Mendelsohn BA, Cope H, Rosenfeld JA, Blackburn PR, Goodenberger ML, Kearney HM, Kennedy J, Scurr I, Szczaluba K, Ploski R, de Saint Martin A, Alembik Y, Piton A, Bruel AL, Thauvin-Robinet C, Strong A, Diderich KEM, Bourgeois D, Dahan K, Vignard V, Bonneau D, Colin E, Barth M, Camby C, Baujat G, Briceño I, Gómez A, Deb W, Conrad S, Besnard T, Bézieau S, Krüger E, Küry S, Stankiewicz P. Stankiewicz-Isidor syndrome: expanding the clinical and molecular phenotype. *Genet Med.* 2022 Jan;24(1):179-191. doi: 10.1016/j.gim.2021.09.005. Epub 2021 Nov 30.

Guest EM, Kairalla JA, Hilden JM, Dreyer ZE, **Carroll AJ**, Heerema NA, Wang CY, Devidas M, Gore L, Salzer WL, Winick NJ, Carroll WL, Raetz EA, Borowitz M, Loh ML, Hunger SP, Brown PA. Outstanding outcomes in infants with KMT2A-germline acute lymphoblastic leukemia treated with chemotherapy alone: results of the Children's Oncology Group AALL0631 trial. *Haematologica.* 2022 Feb 17. doi: 10.3324/haematol.2021.280146. Online ahead of print.

Thomas D, **Soldner JA**, Cropp CD, Beall J. Pharmacy Student Perceptions of a Virtual Pharmacogenomics Activity. *Healthcare (Basel).* 2022 Feb 1;10(2):286. doi: 10.3390/healthcare10020286.

Sadler KV, Rowlands CF, Smith PT, Hartley CL, Bowers NL, Roberts NY, Harris JL, Wallace AJ, Evans DG, **Messiaen LM**, Smith MJ. Re-evaluation of missense variant classifications in NF2. *Hum Mutat.* 2022 Mar 25. doi: 10.1002/humu.24370. Online ahead of print.

Zindl CL, Witte SJ, Laufer VA, **Gao M, Yue Z**, Janowski KM, Cai B, Frey BF, Silberberger DJ, Harbour SN, Singer JR, Turner H, Lund FE, Vallance BA, Rosenberg AF, Schoeb TR, **Chen JY**, Hatton RD, Weaver CT. A nonredundant role for T cell-derived interleukin 22 in antibacterial defense of colonic crypts. *Immunity.* 2022 Mar 8;55(3):494-511.e11. doi: 10.1016/j.immuni.2022.02.003.

Palacka P, Gvozdjaková A, Rausová Z, Kucharská J, Slobovský J, Obertová J, Furka D, Furka S, **Singh KK**, Sumbalová Z. Platelet Mitochondrial Bioenergetics Reprogramming in Patients with Urothelial Carcinoma. *Int J Mol Sci.* 2021 Dec 30;23(1):388. doi: 10.3390/ijms23010388.

Song CJ, Li Z, Ahmed UKB, Bland SJ, Yashchenko A, Liu S, Aloria EJ, Lever JM, Gonzalez NM, Bickel MA, Giles CB, Georgescu C, Wren JD, Lang ML, Benveniste EN, Harrington LE, Tsiokas L, George JF, Jones KL, **Crossman DK**, Agarwal A, Mrug M, Yoder BK, Hopp K, Zimmerman KA. A Comprehensive Immune Cell Atlas of Cystic Kidney Disease Reveals the Involvement of Adaptive Immune Cells in Injury-Mediated Cyst Progression in Mice. *J Am Soc Nephrol.* 2022 Apr;33(4):747-768. doi:

10.1681/ASN.2021030278. Epub 2022 Feb 2.

Schalk A, Cousin MA, Dsouza NR, Challman TD, Wain KE, Powis Z, Minks K, Trimouille A, Lasseaux E, Lacombe D, Angelini C, Michaud V, Van-Gils J, Spataro N, Ruiz A, Gabau E, Stolerman E, Washington C, Louie R, Lanpher BC, Kemppainen JL, Innes M, Kooy F, Meuwissen M, Goldenberg A, Lecoquierre F, Vera G, Diderich KEM, Sheidley B, El Achkar CM, Park M, Hamdan FF, Michaud JL, Lewis AJ, Zweier C, Reis A, Wagner M, Weigand H, Journel H, Keren B, Passemard S, Mignot C, van Gassen K, Brilstra EH, Itzikowitz G, O'Heir E, Allen J, Donald KA, **Korf BR**, Skelton T, Thompson M, **Robin NH**, Rudy NL, Dobyns WB, Foss K, Zarate YA, Bosanko KA, Alembik Y, Durand B, Tran Mau-Them F, Ranza E, Blanc X, Antonarakis SE, McWalter K, Torti E, Millan F, Dameron A, Tokita M, Zimmermann MT, Klee EW, Piton A, Gerard B. De novo coding variants in the AGO1 gene cause a neurodevelopmental disorder with intellectual disability. *J Med Genet*. 2021 Dec 15:jmedgenet-2021-107751. doi: 10.1136/jmedgenet-2021-107751. Online ahead of print.

Angelozzi M, Karvande A, Molin AN, Ritter AL, Leonard JMM, Savatt JM, Douglass K, Myers SM, Grippa M, Tolchin D, Zackai E, Donoghue S, **Hurst ACE, Descartes M**, Smith K, Velasco D, Schmanski A, Crunk A, Tokita MJ, de Lange IM, van Gassen K, Robinson H, Guegan K, Suri M, Patel C, Bournez M, Faivre L, Tran-Mau-Them F, Baker J, Fabie N, Weaver K, Shillington A, Hopkin RJ, Barge-Schaapveld DQCM, Ruivenkamp CA, Bökenkamp R, Vergano S, Seco Moro MN, Díaz de Bustamante A, Misra VK, Kennelly K, Rogers C, Friedman J, Wigby KM, Lenberg J, Graziano C, Ahrens-Nicklas RC, Lefebvre V. Consolidation of the clinical and genetic definition of a SOX4-related neurodevelopmental syndrome. *J Med Genet*. 2022 Mar 1:jmedgenet-2021-108375. doi: 10.1136/jmedgenet-2021-108375. Online ahead of print.

Li G, Holly T, Kelly DR, Reddy V, **Mikhail FM, Carroll AJ**, Kutny MA. Therapy-related Myeloid Neoplasms in Children: A Single-institute Study. *J Pediatr Hematol Oncol*. 2022 Jan 1;44(1):e109-e113. doi: 10.1097/MPH.0000000000002097.

East KM, Cochran ME, Kelley WV, Greve V, Finnila CR, Coleman T, Jennings M, Alexander L, Rahn EJ, Danila MI, Barsh G, **Korf B**, Cooper G. Education and Training of Non-Genetics Providers on the Return of Genome Sequencing Results in a NICU Setting. *J Pers Med*. 2022 Mar 5;12(3):405. doi: 10.3390/jpm12030405.

Patel SB, Nemkov T, Stefanoni D, Benavides GA, Bassal MA, Crown BL, Watkins VR, Camacho V, Kuznetsova V, Hoang AT, Tenen DE, Wolock SL, Park J, Ying L, Yue Z, **Chen JY**, Yang H, Tenen DG, Ferrell PB, Lu R, Darley-Usmar V, D'Alessandro A, Bhatia R, Welner RS. Metabolic alterations mediated by STAT3 promotes drug persistence in CML. *Leukemia*. 2021 Dec;35(12):3371-3382. doi: 10.1038/s41375-021-01315-0. Epub 2021 Jun 12.

Zhang Z, Luo L, Xing C, Chen Y, Xu P, Li M, Zeng L, Li C, Ghosh S, Della Manna D, Townes T, Britt WJ, Wajapeyee N, Sleckman BP, **Chong Z**, Leavenworth JW, Yang ES. RNF2 ablation reprograms the tumor-immune microenvironment and stimulates durable NK and CD4(+) T-cell-dependent antitumor immunity. *Nat Cancer*. 2021 Oct;2(10):1018-1038. doi: 10.1038/s43018-021-00263-z. Epub 2021 Oct 22.

Arora I, Sharma M, Li S, **Crowley M, Crossman DK**, Li Y, Tollefsbol TO. An integrated analysis of the effects of maternal broccoli sprouts exposure on transcriptome and methylome in prevention of offspring mammary cancer. PLoS One. 2022 Mar 9;17(3):e0264858. doi: 10.1371/journal.pone.0264858. eCollection 2022.

Kahan SM, Bakshi RK, Ingram JT, Hendrickson RC, Lefkowitz EJ, **Crossman DK**, Harrington LE, Weaver CT, Zajac AJ. Intrinsic IL-2 production by effector CD8 T cells affects IL-2 signaling and promotes fate decisions, stemness, and protection. Sci Immunol. 2022 Feb 11;7(68):eabl6322. doi: 10.1126/scimmunol.abl6322. Epub 2022 Feb 11.

Yang C, Harafuji N, O'Connor AK, **Kesterson RA**, Watts JA, Majmundar AJ, Braun DA, Lek M, Laricchia KM, Fathy HM, Mane S, Shril S, Hildebrandt F, Guay-Woodford LM. Cystin genetic variants cause autosomal recessive polycystic kidney disease associated with altered Myc expression. Sci Rep. 2021 Sep 14;11(1):18274. doi: 10.1038/s41598-021-97046-4.

Wilson AC, Chiles J, Ashish S, Chanda D, Kumar PL, Mobley JA, Neptune ER, Thannickal VJ, **McDonald MN**. Integrated bioinformatics analysis identifies established and novel TGF β 1-regulated genes modulated by anti-fibrotic drugs. Sci Rep. 2022 Feb 23;12(1):3080. doi: 10.1038/s41598-022-07151-1

Recent Faculty Grant Awards

PI: Michael Crowley

Sponsor: NIH-S10

Title: NovaSeq 6000 Sequencing System

Amount: \$945,108

Dates: 3-1-2022 to 2-28-23

PI: Bruce Korf

Sponsor: Children's Tumor Foundation

Title: NFCTC Data Management Support

Amount: \$50,936

Dates: 2-1-2022 to 1-31-23

PI: Bruce Korf

Sponsor: NF2 Biosolutions

Title: NF2 BioSolutions Blood and Tumor Banking Services for the

Neurofibromatosis Clinical Trials Consortium (NFCTC)

Amount: \$15,000

Dates: 2-26-2022 to 2-25-2023

PI: Robert Kesterson

Sponsor: Gilbert Family Foundation

Title: NF1 Gene Rescue in Translational Animal Models

Amount: \$400,000

Dates: 1-1-2022 to 12-31-22

PI: Keshav Singh

Sponsor: NIH-R21

Title: mtDNA Depleter Mouse for Decoding Mitochondrial Regulation of Diverse Organs

Amount: \$200,475

Dates: 3-15-2022 to 2-28-23

PI: Zhangli Su

Sponsor: NIH-K99

Title: The Role of Small RNA Modifications in Glioma

Amount: \$110,646

Dates: 3-11-2022 to 2-28-23

PI: Deeann Wallis

Sponsor: Gilbert Family Foundation

Title: Targeted Delivery of Full Length NF1 cDNA using Non-viral Vectors

Amount: \$386,697

Dates: 1-1-2022 to 12-31-22

PI: Deeann Wallis

Sponsor: Gilbert Family Foundation

Title: RAFT-Polymerized Nanoparticles for Delivery of NF1 cDNA to Schwann Cells

Amount: \$28,480

Dates: 10-1-2021 to 9-30-22

PI: Deeann Wallis

Sponsor: Gilbert Family Foundation

Title: Exon Skipping to Treat NF1

Amount: \$347,082

Dates: 2-1-2022 to 1-31-23

Department Acknowledgements

Dan Sharer, Ph.D., professor, assumed the role of Interim Director of Genetics Research in January 2022.

Anna Hurst, M.D., assistant professor, was named associate editor of the American Journal of Medical Genetics.

Bruce Korf, M.D., Ph.D., professor, was elected to the American Association of Physicians.

Yulong Fu, Ph.D., assistant professor, named interim director of the Medical Genomics Laboratory.

Steve Doran, named space and facilities manager.

Thank you to **Leigh Ann Withers** and **Shelia Coleman** for all of their work on faculty recruitment recently.

Congratulations to the **UAB Laboratory Genetics and Genomics (LGG) ABMGG Fellowship** on being accredited by the ACGME.

Welcome to the Department

New Staff	New Post-Docs	New Students
Collin Ainslie	Shruti Kapoor	Logan Brewer
Elizabeth Michelle Bird	Pradeep Kumar	Amy Lamore
Rose Missildine	Sarmad Mehmood	Violet Morin
Annah Bochaberi Oigo	Nongyao Nonpanya	Danya Midani
Isaac Andres Segura Rueda		Holland Powell
Xiaoxia Zhang		Brayden Rohr
		Kylee Spencer

Upcoming Events

Save the Date: Department Faculty and Staff Retreat - Wednesday, May 25



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Editor: [Heather Watts](#)

If you have content you would like included in the quarterly newsletter,
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