



Flow Cytometry Publications List

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Bulletin 6741



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online article.

Cancer

Catenacci DV et al. (2015).

Acquisition of portal venous circulating tumor cells from patients with pancreaticobiliary cancers by endoscopic ultrasound.



Gastroenterology 149, 1,794–1,803.

Frame FM et al. (2016).

Harvesting human prostate tissue material and culturing primary prostate epithelial cells.



Methods Mol Biol 1443, 181–201.

Hernandez JR et al. (2015).

Alternative CD44 splicing identifies epithelial prostate cancer cells from the mesenchymal counterparts.



Med Oncol 32, 159.

Jin HJ et al. (2016).

Identification and validation of regulatory SNPs that modulate transcription factor chromatin binding and gene expression in prostate cancer.



Oncotarget 7, 54,616–54,626.

Kawahara T et al. (2016).

Silodosin inhibits prostate cancer cell growth via ELK1 inactivation and enhances the cytotoxic activity of gemcitabine.



Prostate 76, 744–756.

Lacotte S et al. (2016).

Impact of myeloid-derived suppressor cell on Kupffer cells from mouse livers with hepatocellular carcinoma.



Oncoimmunology 5, e1234565.

Li Y et al. (2015).

P16INK4A is required for cisplatin resistance in cervical carcinoma SiHa cells.



Oncol Lett 9, 1,104–1,108.



Cancer *continued*

Peirs S et al. (2014).

ABT-199 mediated inhibition of BCL-2 as a novel therapeutic strategy in T-cell acute lymphoblastic leukemia.

Blood 124, 3,738–3,747.



Rae DT et al. (2015).

A novel retroviral mutagenesis screen identifies prognostic genes in RUNX1 mediated myeloid leukemogenesis.

Oncotarget 6, 30,664–30,674.



Rihani A et al. (2015).

Inhibition of CDK4/6 as a novel therapeutic option for neuroblastoma.

Cancer Cell Int 15, 76.



Cell Characterization

Bezzerides VJ et al. (2016).

CITED4 induces physiologic hypertrophy and promotes functional recovery after ischemic injury.

JCI Insight 1, e85904.



Blomen VA et al. (2015).

Gene essentiality and synthetic lethality in haploid human cells.

Science 350, 1,092–1,096.



Hervier B et al. (2016).

Involvement of NK cells and NKp30 pathway in antisynthetase syndrome.

J Immunol 197, 1,621–1,630.



Konda N et al. (2016).

Truncated EphA2 likely potentiates cell adhesion via integrins as well as infiltration and/or lodgment of a monocyte/macrophage cell line in the red pulp and marginal zone of the mouse spleen, where ephrin-A1 is prominently expressed in the vasculature.

Histochem Cell Biol [published ahead of print September 24, 2016]. Accessed

February 14, 2017.



Lother A et al. (2016).

Deoxycorticosterone acetate/salt-induced cardiac but not renal injury is mediated by endothelial mineralocorticoid receptors independently from blood pressure.

Hypertension 67, 130–138.



Nührenberg TG et al. (2015).

Cardiac myocyte de novo DNA methyltransferases 3a/3b are dispensable for cardiac function and remodeling after chronic pressure overload in mice.

PLoS One 10, e0131019.





Cell Characterization *continued*

Regn M et al. (2016).

Peptidase inhibitor 16 is a membrane-tethered regulator of chemerin processing in the myocardium.

J Mol Cell Cardiol 99, 57–64.



Turqueti-Neves A et al. (2015).

The extracellular domains of IgG1 and T cell-derived IL-4/IL-13 are critical for the polyclonal memory IgE response in vivo.

PLoS Biol 13, e1002290.



Cell Development

Džinić T et al. (2016).

Oxygen and differentiation status modulate the effect of X-ray irradiation on physiology and mitochondrial proteome of human neuroblastoma cells.

Arch Physiol Biochem 122, 257–265.



Gilsbach R et al. (2014).

Dynamic DNA methylation orchestrates cardiomyocyte development, maturation and disease.

Nat Commun 5, 5,288.



Püngel S et al. (2015).

Reconciling pillars of transient gene expression: From DNA prep via media, reagent and cell line development to holistic process optimization.

BMC Proc 9, P18.



Saeki N et al. (2015).

EphA2 promotes cell adhesion and spreading of monocyte and monocyte/macrophage cell lines on integrin ligand-coated surfaces.

Cell Adh Migr 9, 469–482.



Cell Differentiation

Afzal E et al. (2013).

Nanolipodendrosome-loaded glatiramer acetate and myogenic differentiation 1 as augmentation therapeutic strategy approaches in muscular dystrophy.

Int J Nanomedicine 8, 2,943–2,960.



Freedman BD et al. (2013).

Adrenocortical zonation results from lineage conversion of differentiated zona glomerulosa cells.

Dev Cell 26, 666–673.





Cell Differentiation *continued*

Nam S et al. (2016).

Interferon regulatory factor 4 (IRF4) controls myeloid-derived suppressor cell (MDSC) differentiation and function.



J Leukoc Biol 100, 1,273–1,284.

Onder L et al. (2015).

Alternative NF- κ B signaling regulates mTEC differentiation from podoplanin-expressing precursors in the cortico-medullary junction.



Eur J Immunol 45, 2,218–2,231.

Sécca C et al. (2016).

IRF2BP2 transcriptional repressor restrains naive CD4 T cell activation and clonal expansion induced by TCR triggering.



J Leukoc Biol 100, 1,081–1,091.

Cell Identification

Carulli G et al. (2015).

Combination of CD157 and FLAER to detect peripheral blood eosinophils by multiparameter flow cytometry.



J Clin Exp Hematop 55, 55–60.

Cell Sorting

Riddell A et al. (2015).

Rmax: A systematic approach to evaluate instrument sort performance using center stream catch.



Methods 82, 64–73.

Epigenetics

Preissl S et al. (2015).

Deciphering the epigenetic code of cardiac myocyte transcription.



Circ Res 117, 413–423.

Tang WW et al. (2015).

A unique gene regulatory network resets the human germline epigenome for development.



Cell 161, 1,453–1,467.



Immunology

Gil-Cruz C et al. (2016).

Fibroblastic reticular cells regulate intestinal inflammation via IL-15-mediated control of group 1 ILCs.

Nat Immunol 17, 1,388–1,396.



Imhof BA et al. (2016).

CCN1/CYR61-mediated meticulous patrolling by Ly6Clow monocytes fuels vascular inflammation.

Proc Natl Acad Sci USA 113, E4,847–E4,856.



Menendez CM et al. (2016).

Resident T cells are unable to control herpes simplex virus-1 activity in the brain ependymal region during latency.

J Immunol 197, 1,262–1,275.



Microbial Analysis

Bidlingmaier S et al. (2016).

Proteome-wide identification of novel ceramide-binding proteins by yeast surface cDNA display and deep sequencing.

Mol Cell Proteomics 15, 1,232–1,245.



Eshaghi M et al. (2016).

Brighter fluorescent derivatives of UTI89 utilizing a monomeric vGFP.

Pathogens 5, E3.



Guizou S et al. (2016).

A part toolbox to tune genetic expression in *Bacillus subtilis*.

Nucleic Acids Res 44, 7,495–7,508.



Guo X et al. (2015).

Innate lymphoid cells control early colonization resistance against intestinal pathogens through ID2-dependent regulation of the microbiota.

Immunity 42, 731–743.



Lander N et al. (2015).

CRISPR/Cas9-induced disruption of paraflagellar rod protein 1 and 2 genes in *Trypanosoma cruzi* reveals their role in flagellar attachment.

MBio 6, e01012– e01015.



Liu Y et al. (2016).

Immune activation of the host cell induces drug tolerance in *Mycobacterium tuberculosis* both in vitro and in vivo.

J Exp Med 213, 809–825.



Moore SJ et al. (2016).

EcoFlex: A multifunctional MoClo kit for *E. coli* synthetic biology.

ACS Synth Biol 5, 1,059–1,069.





Protein Analysis

Eshaghi M et al. (2015).

Rational structure-based design of bright GFP-based complexes with tunable dimerization.

Angew Chem Int Ed Engl 54, 13,952–13,956.



Martínez-Limón A et al. (2016).

Recognition of enzymes lacking bound cofactor by protein quality control.

Proc Natl Acad Sci USA 113, 12,156–12,161.



Whiten DR et al. (2016).

Rapid flow cytometric measurement of protein inclusions and nuclear trafficking.

Sci Rep 6, 31138.



Stem Cells

Bertacchi M et al. (2015).

Activin/nodal signaling supports retinal progenitor specification in a narrow time window during pluripotent stem cell neuralization.

Stem Cell Reports 5, 532–545.



Chal J et al. (2015).

Differentiation of pluripotent stem cells to muscle fiber to model Duchenne muscular dystrophy.

Nat Biotechnol 33, 962–969.



Irie N and Surani MA (2017).

Efficient induction and isolation of human primordial germ cell-like cells from competent human pluripotent stem cells.

Methods Mol Biol 1463, 217–226.



Murakami K et al. (2016).

NANOG alone induces germ cells in primed epiblast in vitro by activation of enhancers.

Nature 529, 403–407.



Pacini S et al. (2016).

Mesangiogenic progenitor cells derived from one novel CD64(bright) CD31(bright)CD14(neg) population in human adult bone marrow.

Stem Cells Dev 25, 661–673.



Quintanilla RH Jr et al. (2016).

Kinetic measurement and real time visualization of somatic reprogramming.

J Vis Exp, issue 113 [video].





Virology

Browning DL et al. (2016).

Insulated foamy viral vectors.

Hum Gene Ther 27, 255–266.



Ramanujam D et al. (2016).

Viral vector-based targeting of miR-21 in cardiac nonmyocyte cells reduces pathologic remodeling of the heart.

Mol Ther 24, 1,939–1,948.



Yadav J et al. (2014).

Opportunistic infections and complications in human immunodeficiency virus-1-infected children: Correlation with immune status.

Sultan Qaboos Univ Med J 14, e513–e521.



Flow Cytometry Posters

Lakshmipathy U et al.

Real time visualization and kinetic measurement of somatic reprogramming.

Presented at: International Society for Stem Cell Research; June 24–27, 2015;
Stockholm, Sweden.



Litterst L et al.

Rapid and ultra-sensitive single-cell transcript profiling with Droplet Digital PCR (ddPCR™): Application to neuronal differentiation.

Presented at: Neuroscience 2014; November 15–19, 2014; Washington, D.C.



Riddell A.

Using a spectra physics Tsunami infrared pulsed laser and second harmonic generation on a Beckman Coulter MoFlo Legacy sorter: A tunable light source.

Presented at: 30th Congress of the International Society for Advancement of Cytometry; June 26–30, 2015; Glasgow, Scotland. Abstract 421/B290, 248.



Riddell A et al.

Sorting and maintaining haploid stem cells using a Beckman Coulter MoFlo Legacy and a Bio-Rad S3 by light scatter.

Presented at: 30th Congress of the International Society for Advancement of Cytometry; June 26–30, 2015; Glasgow, Scotland. Abstract 420/B289, 247–248.



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