

## **Tiziana Bonaldi, PhD**

### **PERSONAL INFORMATION**

Family name, first name: Bonaldi, Tiziana  
Researcher unique identifier(s) 0000-0003-3556-1265 (ORCID ID)  
Date of birth: 04 July 1973  
Nationality: Italian  
URL for web site: <https://www.ieo.it/it/ricerca/People/Ricercatori/Bonaldi-Tiziana/>

### **EDUCATION**

**2003** PhD award of the Open University of London, program held at DiBit, University Vita Salute, San Raffaele Hospital (HSR), Milan, IT (PhD supervisors: Internal Prof. M.E. Bianchi; external Prof. B. Turner (University of Birmingham, UK))  
**1997** B. Sc. (Laurea) at the University of Milan, Italy- *Summa cum Laudae*

### **CURRENT POSITIONS**

**2015** Tenured Group Leader at the Department of Experimental Oncology, European Institute of Oncology, IEO, IT  
**2014** MIUR (Italian Ministry of Schools, University and Research) Habilitation as associate Professor of Biochemistry, Molecular Biology and Applied Biology  
**2015** Member the Scientific Directors Committee of SIBBM (Italian Society of Biophysics and molecular Biology, [http://www.sibbm.org/home .php](http://www.sibbm.org/home.php))  
**2008** Faculty member of European School of Molecular Medicine (SEMM) PhD Program, Milan

### **PREVIOUS POSITIONS**

**2008 - 2014** Junior Group Leader (tenure-track) at the Department of Experimental Oncology, European Institute of Oncology, IT  
**2006 - 2008** Post-doctoral fellow at Department of Proteomics and Signal Transduction the Max Plank Institute of Biochemistry, Martinsried, DE (Head: Prof. Matthias Mann)  
**2003 - 2005** EMBO Long-Term Post-doctoral fellow in the MS and Chromatin Composition Unit (Head: Prof. Axel Imhof) at Ludwig Maximilians University (LMU), Munich, DE  
**2002 - 2003** Visiting scientist in the Mass Spectrometry Group, at the European Molecular Biology Lab (EMBL), DE (Head: Dr. Matthias Wilm)

### **FELLOWSHIPS, AWARDS & HONOURS**

**2010** International Inner Wheel for Women 2010, for scientific career  
**2007** The Giovanni Armenise Harvard Foundation career development Award  
**2003 - 2005** EMBO Long-Term Post-Doctoral Fellowship at the LMU, DE  
**2000** EMBO Short-Term Fellowship. DiBit-San Raffaele Hospital, IT  
**1999 - 2003** 1<sup>st</sup> ranked Doctoral Fellowship offered by University of Milan, IT

### **CAREER BREAKS**

Five months Maternity leave, for the birth of my first child, Fernanda Stabilini  
Five Months Maternity leave, for the birth of my second child, Pietro Stabilini

## **ACADEMIC ACTIVITY**

### **Supervision of graduate students and postdoctoral fellows**

**2008 - 2017** As PI, I supervised 10 post-docs/ 4 PhD students/ 7 Master students (4 Italians) at the Department of Experimental Oncology–IEO, Milan-Bicocca University, University' degli Studi di Milano, University of Parma.

I have hosted in my Group 5 foreigner visiting student for short term period (1 to 4 months) for training in MS-proteomics.

**2006 - 2008** As a postdoc, I supervised 1 PhD student in M. Mann's group at the Max Planck of Biochemistry in Martinsried, DE

### **Teaching activity**

**2015** Lecturer for monographic short course (4 hours) within the course of Clinical Biochemistry at the University of Piemonte Orientale, Novara, IT

**2014** Lecturer at PhD course "Mass-spectrometry and Proteomics in cell biology research" at University of Padua (Padua)

**2010 and 2017-** Lecturer PhD course in "Mass Spectrometry-based proteomics and its applications in biology", NNF Center for Protein Research, Copenhagen

**2010** Lecturer at the PhD course "Mass-spectrometry and Proteomics in cell biology research" at CNR Institute of Neuroscience, Milano

**2010** Lecturer at the PhD course in Epigenetics at Dept of Genetics University of Milano

**Since 2008** Assistant Professor and coordinator of the course "Proteomics" for the School of Molecular Medicine (SEMM);

### **Participation to Examination/Evaluation Board**

**2017** Istituto di Genetica Molecolare (IGM) of the CNR: Competition for scientist (ricercatore) appointment

**2016** Biochemistry Department of the Technical University (TU) of Munich: Evaluation of candidate for the promotion to the position of "adjunct professor"

**2016** Department of Veterinary of the University of Milano: Competition for post-doc position appointment

**2016** Koc University (KU), Istanbul (TK): evaluation of candidate for promotion as Associate Professor

### **Lecturer and Trainer at International Conferences**

**2015** 9<sup>th</sup> European Summer School on Advanced Proteomics, Brixen (IT)

**2010** HUPO2010 Education and Training day; Sydney (AU)

**2007** Practical Course "MS-based Proteomics and Quantitative Proteomics" at the Max Planck Institute of Biochemistry, Martinsried (DE) (2007);

## **EDITORIAL ACTIVITY**

**Editorial Board Member:** Journal of Integrated –OMICS, Frontiers in Molecular and Cellular Oncology, Advances in Biology, Molecular and Cellular Oncology; International Journal of Proteomics, Proteomes.

**Ad hoc peer-reviewer:** Nature Protocols, *Biochimica and Biophysica Acta*, Proteomics, Journal of Proteome Research, Journal of Cell Science, Journal of Proteomics, Trends in Analytical Chemistry, Expert Reviews in Proteomics, Briefings in Functional Genomics, BMC Cancer, Trends in Biotechnology, Nature Communications, NAR, Journal of Molecular Cell Biology, Molecular and Cellular Proteomics, Oncogene, Scientific Reports, Epigenetic & Chromatin, Oncotarget, Plos One

**Grant Reviewer:** AICR, FEBS, Research Foundation Flanders, CRUK, Wellcome Trust, Agencie National Recherche (ANR), Cancer Research HK, ERC-stating Grant (external expert reviewer), German-Israeli (DKFZ-MOST) cooperation in cancer research.

**SCIENTIFIC MEETINGS ORGANIZER and co-organizer**

**2017** SIBBM Annual Meeting, Milano: organizer and chair of Session

**2016** 2<sup>nd</sup> EMBO | FEBS Lecture Course on Nuclear Proteomics, Crete, GRI International Conference Proteomics & Bioinformatics, Rome- IT| 4<sup>th</sup> and 5<sup>th</sup> MS-proteomics Days, Milan Area, Italy| SIBBM meeting, Naples, IT

**2015** International Conference Proteomics & Bioinformatics, Valencia, ESI 2<sup>nd</sup> and 3<sup>rd</sup> MS-Proteomics Day, Milano Area, IT

**2014** 1<sup>st</sup> MS-Proteomics Day, IFOM-IEO Campus, Milan, IT| EMBO- FEBS Lecture Course on Nuclear Proteomics, Kos Island, GRI International Conference Proteomics & Bioinformatics, Chicago, USA| EPIGEN-MiChroNetwork Chromatin Seminar, Dept. Experimental Oncology, IEO

**2011** Epigenetic Symposium, IEO, Milan, IT

**2009** Italian Proteomics Association 4<sup>th</sup> National Conference, Milano Bicocca University

**INVITED PRESENTATIONS**

**2017:** Budapest Breast Think Tank EACR Meeting, Budapest HUN (*invited speaker*)| Gordon Research Conference “Cancer Genetics & Epigenetics”, Lucca IT (*invited speaker*)| 5th ICAP Conference, Lisbon POR (*invited speaker*)| EU-LIFE Scientific Workshop "Principles of Homeostasis", Berlin DE (*invited speaker*)| EMBO course “Non-coding RNAs”, Heidelberg DE (*lecturer*)| HUPO2017, Dublin IRE (*invited speaker*)| SIB Annual Meeting, Caserta IT

**2016:** Belgian Proteomics Association (BePac) Meeting, Ghent- BE (*Keynote lecturer*)| FEBS- Nuclear Proteomics Summer School, Heraklion- GR (*Invited speaker and chairman*)| EMBL–WELLCOME GENOME CAMPUS CONFERENCE, Heidelberg- DE (*Invited speaker and chairman*)| Giovanni Armenise-Harvard 16<sup>th</sup> Symposium- Gubbio– IT (*Invited speaker*)| SIBBM Meeting- Naples- IT (*Organizer and chairman*)| EPIGEN-National Meeting, Rome- IT (*Invited speaker*)| Game of Epigenomics –Dubrovnik- CRO (*Invited speaker*)| Lecture Babraham Institute- Cambridge –UK (*Invited speaker*)| Lecture at the CRUK Beatson Institute- Glasgow-UK (*Invited speaker*)

**2015:** International Conference Proteomics & Bioinformatics, Valencia- SP (*speaker & chairman*)| Hallmarks of Cancer: focus on RNA, Paris- FR (*selected talk*)| GAHF 15<sup>th</sup> symposium, Challenges to Modern Medicine, Rome - IT (*invited speaker*)

**2014:** EPIGEN Annual Meeting, Rome- IT | 2<sup>nd</sup> Epigenetic Symposium, Oslo- NO| SIBBM Trento-IT| FISV 2014, Pisa- IT (*invited speaker and chairman*)| 4D Cell Symposium, Utrecht (HOL)

**2013:** Proteomic Forum 2013, Berlin- DE| 7<sup>th</sup> ITPA Annual Congress, Padova- IT

**2012:** SIBBM 2012 Frontiers in molecular biology, Palermo- IT| 11<sup>th</sup> National Meeting of Medicinal Chemistry, Palermo- IT| 9<sup>th</sup> Siena Meeting from Genome to Proteome, IT| FISV Conference, Rome- IT

**2011:** II International Congress on Analytical Proteomics, Ourense -SP

**2010:** Marie-Curie Network INTEGER “Chromatin Proteomics” meeting, Munich –DE| Italian Proteomics Association 5th Annual National Conference, Firenze- IT| HUPO 9<sup>th</sup> Annual World Conference 2010, Sydney-AUS (*lecturer and invited speaker*)

**2009:** 11<sup>th</sup> International Congress on Amino Acids, Peptides and Proteins, Vienna- AT| Open Symposium: “20 years of Electrospray MS for large molecules”, Munich- DE| Italian Proteomics Association 4<sup>th</sup> Annual National Conference, Milano- IT (*co-organizer*)

## **CITATION METRICS (WEB OF SCIENCE March 2018)**

*Number of publications:* 63

*h-index:* 27

## **PEER-REVIEWED PUBLICATIONS (LAST 12 YEAR)**

### **2018**

1. Noberini R, Osti D, Miccolo C, Richichi C, Lupia M, Corleone G, Hong SP, Colombo P, Pollo B, Fornasari L, Pruneri G, Magnani L, Cavallaro U, Chiocca S, Minucci S, Pelicci G, **Bonaldi T**. Extensive and systematic rewiring of histone post-translational modifications in cancer model systems. *Nucleic Acids Res.* 2018 Mar 29. doi: 10.1093/nar/gky224.
2. Audano M, Pedretti S, Cermenati G, Brioschi E, Diaferia GR, Ghisletti S, Cuomo A, **Bonaldi T**, Salerno F, Mora M, Grigore L, Garlaschelli K, Baragetti A, Bonacina F, Catapano AL, Norata GD, Crestani M, Caruso D, Saez E, De Fabiani E, Mitro N. Zc3h10 is a novel mitochondrial regulator. *EMBO Rep.* 2018 Mar 5. pii: e45531. doi: 10.15252/embr.201745531.
3. Caprara G, Prosperini E, Piccolo V, Sigismondo G, Melacarne A, Cuomo A, Boothby M, Rescigno M, **Bonaldi T**, Natoli G. PARP14 Controls the Nuclear Accumulation of a Subset of Type I IFN-Inducible Proteins. *J Immunol.* 2018 Apr 1;200(7):2439-2454. doi: 10.4049/jimmunol.1701117.
4. Fellows R, Denizot J, Stellato C, Cuomo A, Jain P, Stoyanova E, Balázs S, Hajnády Z, Liebert A, Kazakevych J, Blackburn H, Corrêa RO, Fachi JL, Sato FT, Ribeiro WR, Ferreira CM, Perée H, Spagnuolo M, Mattiuz R, Matolcsi C, Guedes J, Clark J, Veldhoen M, **Bonaldi T**, Vinolo MAR, Varga-Weisz P. Microbiota derived short chain fatty acids promote histone crotonylation in the colon through histone deacetylases. *Nat Commun.* 2018 Jan 9;9(1):105. DOI: 10.1038/s41467-017-02651-5.

### **2017**

1. Hoffmann G, Samel-Pommerencke A, Weber J, Cuomo A, **Bonaldi T**, Ehrenhofer-Murray AE. A role for CENP-A/Cse4 phosphorylation on serine 33 in deposition at the centromere. *FEMS Yeast Res.* 2017 Dec 20. doi: 10.1093/femsyr/fox094.
2. Soldi M, Mari T, Nicosia L, Musiani D, Sigismondo G, Cuomo A, Pavesi G, **Bonaldi T**. Chromatin proteomics reveals novel combinatorial histone modification signatures that mark distinct subpopulations of macrophage enhancers *Nucleic Acids Research*, 2017 Dec 1;45(21):12195-12213. DOI: 10.1093/nar/gkx821
3. Chen YK, **Bonaldi T**, Cuomo A, Del Rosario JR, Hosfield DJ, Kanouni T, Kao S, Lai C, Lobo NA, Matuszkiewicz j, McGeehan A, O'Connell SM, Shi L, Stafford JA, Stansfield RK, Veal JM, Weiss MS, Yuen NY, and Wallace MB. Design of KDM4 Inhibitors with Antiproliferative Effects in Cancer Models. *ACS Med. Chem. Lett.*, **2017**, 8 (8), pp 869–874. DOI: 10.1021/acsmchemlett.7b00220.
4. Noberini R, Longuespée R, Richichi C, Pruneri G, Kriegsmann M, Pelicci G., **Bonaldi T**. PAT-H-MS coupled with laser microdissection to study histone post-translational modifications in selected cell populations from pathology samples. *Clin Epigenetics.* 2017 Jul 11;9:69. doi: 10.1186/s13148-017-0369-8. eCollection **2017**. PubMed PMID: 28702092; PubMed Central PMCID: PMC5504751.
5. Caldieri G, Barbieri E, Nappo G, Raimondi A, Bonora M, Conte A, Verhoef LGGC, Confalonieri S, Malabarba MG, Bianchi F, Cuomo A, **Bonaldi T**, Martini E, Mazza D, Pinton P, Tacchetti C, Polo S, Di Fiore PP, Sigismund S. Reticulon 3-dependent ER-PM contact sites control EGFR nonclathrin endocytosis. *Science.* **2017** May 12;356(6338):617-

624. doi: 10.1126/science.aah6152. PubMed PMID: 28495747; PubMed Central PMCID: PMC5432029.

6. Curina A, Termanini A, Barozzi I, Prosperini E, Simonatto M, Polletti S, Silvola A, Soldi M, Austenaa L, **Bonaldi T**, Ghisletti S, Natoli G. High constitutive activity of a broad panel of housekeeping and tissue-specific cis-regulatory elements depends on a subset of ETS proteins. *Genes Dev.* 2017 Feb 15;31(4):399-412. doi: 10.1101/gad.293134.116. Epub 2017 Mar 8. PubMed PMID: 28275002; PubMed Central PMCID: PMC5358759.

## **2016**

7. Noberini R, Pruneri G, Minucci S, **Bonaldi T**. Mass-spectrometry analysis of histone post-translational modifications in pathology tissue using the PAT-H-MS approach. *Data Brief.* 2016 Feb 16;7:188-94. doi: 10.1016/j.dib.2016.02.028. eCollection 2016 Jun. PubMed PMID: 27408908; PubMed Central PMCID: PMC4927966.
8. Soldi M, Cuomo A, **Bonaldi T**. Quantitative assessment of chemical artefacts produced by propionylation of histones prior to mass spectrometry analysis. *Proteomics.* 2016 Jul;16(14):1952-4. doi: 10.1002/pmic.201600173. PubMed PMID: 27373704
9. **Bonaldi T**, Mihailovich M. Spatiotemporal plasticity of miRNAs functions: The miR-17-92 case. *Mol Cell Oncol.* 2016 Apr 15;3(3):e1156216. doi: 10.1080/23723556.2016.1156216. eCollection 2016 May. PubMed PMID: 27314099; PubMed Central PMCID: PMC4909434
10. Pozzi C, Cuomo A, Spadoni I, Magni E, Silvola A, Conte A, Sigismund S, Ravenda PS, **Bonaldi T**, Zampino MG, Cancelliere C, Di Fiore PP, Bardelli A, Penna G, Rescigno M. The EGFR-specific antibody cetuximab combined with chemotherapy triggers immunogenic cell death. *Nat Med.* 2016 Jun;22(6):624-31. doi: 10.1038/nm.4078. Epub 2016 May 2. PubMed PMID: 27135741.
11. Mihailovich M, **Bonaldi T**. MYC/miR-17-92 interplay maintains B-lymphoma cell homeostasis. *Cell Cycle.* 2016;15(8):1025-6. doi: 10.1080/15384101.2016.1157977. PubMed PMID: 27097369; PubMed Central PMCID: PMC4889254.
12. Mihailovich M, **Bonaldi T**. MS-analysis of SILAC-labeled MYC-driven B lymphoma cells overexpressing miR-17-19b. *Data Brief.* 2016 Feb 24;7:349-53. doi: 10.1016/j.dib.2016.02.031. eCollection 2016 Jun. PubMed PMID: 26977435; PubMed Central PMCID: PMC4781929.

## **2015**

13. Noberini R, Sigismondo G, **Bonaldi T**. The contribution of mass spectrometry-based proteomics to understanding epigenetics. *Epigenomics.* 2015 Nov 25 [Epub ahead of print] PubMed PMID: 26606673.
14. Mihailovich M, Bremang M, Spadotto V, Musiani D, Vitale E, Varano G, Zambelli F, Mancuso FM, Cairns DA, Pavesi G, Casola S, **Bonaldi T**. miR-17-92 fine-tunes MYC expression and function to ensure optimal B cell lymphoma growth. *Nat Commun.* 2015 Nov 10;6:8725. doi: 10.1038/ncomms9725. PubMed PMID: 26555894.
15. Noberini R, Uggetti A, Pruneri G, Minucci S, **Bonaldi T**. Pathology tissue-quantitative mass spectrometry analysis to profile histone post-translational modification patterns in patient samples. *Mol Cell Proteomics.* 2015 Oct 13. pii: mcp.M115.054510. [Epub ahead of print] PubMed PMID: 26463340.
16. Setti M, Osti D, Richichi C, Ortensi B, Bene MD, Fornasari L, Beznoussenko G, Mironov A, Rappa G, Cuomo A, Faretta M, **Bonaldi T**, Lorico A, Pelicci G. Extracellular vesicle-mediated transfer of CLIC1 protein is a novel mechanism for the regulation of

- glioblastoma growth. *Oncotarget*. **2015** Sep 30. [Epub ahead of print] PubMed PMID: 26429879.
17. Carissimi C, Laudadio I, Cipolletta E, Gioiosa S, Mihailovich M, **Bonaldi T**, Macino G, Fulci V. ARGONAUTE2 cooperates with SWI/SNF complex to determine nucleosome occupancy at human Transcription Start Sites. *Nucleic Acids Res*. **2015** Feb 18;43(3):1498-512. doi: 10.1093/nar/gku1387. Epub 2015 Jan 20. PubMed PMID: 25605800; PubMed Central PMCID: PMC4330357.

#### **2014**

18. Marchesi S, Montani F, Deflorian G, D'Antuono R, Bologna S, Mazzoccoli C, Cuomo A, **Bonaldi T**, Di Fiore PP and Nicassio F "DEPDC1B coordinates de-adhesion events and cell cycle progression at mitosis" *Developmental Cell*, **2014** Nov 24;31(4):420-33.doi: 10.1016/j.devcel.2014.09.009. Epub 2014 Nov 24. PubMed PMID: 25458010;PubMed Central PMCID: PMC4250264.
19. Soldi M, Cuomo A, **Bonaldi T**. Improved bottom-up strategy to efficiently separate hyper-modified histone peptides through ultra HPLC separation on a benchtop Orbitrap instrument. *Proteomics*. **2014** Jul 29. doi: 10.1002/pmic.201400075. [Epub ahead of print] PubMed PMID: 25073962.
20. Soldi M, **Bonaldi T**. The ChroP approach combines ChIP and mass spectrometry to dissect locus-specific proteomic landscapes of chromatin. *J Vis Exp*. **2014** Apr 11;(86). doi: 10.3791/51220. PubMed PMID: 24747196.
21. Colzani M, Noberini R, Romanenghi M, Colella G, Pasi M, Fancelli D, Varasi M, Minucci S, **Bonaldi T**. Quantitative chemical proteomics identifies novel targets of the anti-cancer multi-kinase inhibitor E-3810. *Mol Cell Proteomics*. **2014** Jun;13(6):1495-509. doi: 10.1074/mcp.M113.034173. Epub 2014 Apr 2. PubMed PMID: 24696502; PubMed Central PMCID: PMC4047469.
22. Soldi M, Bremang M, **Bonaldi T**. Biochemical systems approaches for the analysis of histone modification readout. *Biochim Biophys Acta*. **2014** Aug;1839(8):657-68. doi: 10.1016/j.bbagr.2014.03.008. Epub 2014 Mar 27. PubMed PMID: 24681439.
23. Ferrari KJ, Scelfo A, Jammula S, Cuomo A, Barozzi I, Stützer A, Fischle W, **Bonaldi T**, Pasini D. Polycomb-dependent H3K27me1 and H3K27me2 regulate active transcription and enhancer fidelity. *Mol Cell*. **2014** Jan 9;53(1):49-62. doi: 10.1016/j.molcel.2013.10.030. Epub 2013 Nov 27. PubMed PMID: 24289921.

#### **2013**

24. Sigismund S, Algisi V, Nappo G, Conte A, Pascolutti R, Cuomo A, **Bonaldi T**, Argenzio E, Verhoef LG, Maspero E, Bianchi F, Capuani F, Ciliberto A, Polo S, Di Fiore PP. Threshold-controlled ubiquitination of the EGFR directs receptor fate. *EMBO J*. **2013** Jul 31;32(15):2140-57. doi: 10.1038/emboj.2013.149. Epub 2013 Jun 25. PubMed PMID: 23799367; PubMed Central PMCID: PMC3730230.
25. Bremang M, Cuomo A, Agresta AM, Stugiewicz M, Spadotto V, **Bonaldi T**. Mass spectrometry-based identification and characterisation of lysine and arginine methylation in the human proteome. *Mol Biosyst*. **2013** Sep;9(9):2231-47. doi:10.1039/c3mb00009e. PubMed PMID: 23748837.
26. Fragola G, Germain PL, Laise P, Cuomo A, Blasimme A, Gross F, Signaroldi E, Bucci G, Sommer C, Pruneri G, Mazzarol G, **Bonaldi T**, Mostoslavsky G, Casola S, Testa G. Cell reprogramming requires silencing of a core subset of polycomb targets. *PLoS Genet*. **2013**;9(2):e1003292. doi: 10.1371/journal.pgen.1003292. Epub 2013 Feb 28. PubMed PMID: 23468641; PubMed Central PMCID: PMC3585017.

27. Soldi M, Cuomo A, Bremang M, **Bonaldi T**. Mass spectrometry-based proteomics for the analysis of chromatin structure and dynamics. *Int J Mol Sci.* **2013** Mar 6;14(3):5402-31. doi: 10.3390/ijms14035402. PubMed PMID: 23466885; PubMed Central PMCID: PMC3634404.
28. Greiner D, **Bonaldi T**, Eskeland R, Roemer E, Imhof A. Reply to "Chaetocin is a nonspecific inhibitor of histone lysine methyltransferases". *Nat Chem Biol.* **2013** Mar;9(3):137. doi: 10.1038/nchembio.1188. PubMed PMID: 23416388.
29. Vella P, Scelfo A, Jammula S, Chiacchiera F, Williams K, Cuomo A, Roberto A, Christensen J, **Bonaldi T**, Helin K, Pasini D. Tet proteins connect the O-linked N-acetylglucosamine transferase Ogt to chromatin in embryonic stem cells. *Mol Cell.* **2013** Feb 21;49(4):645-56. doi: 10.1016/j.molcel.2012.12.019. Epub 2013 Jan 24. PubMed PMID: 23352454.
30. Soldi M, **Bonaldi T**. The proteomic investigation of chromatin functional domains reveals novel synergisms among distinct heterochromatin components. *Mol Cell Proteomics.* **2013** Mar;12(3):764-80. doi: 10.1074/mcp.M112.024307. Epub 2013 Jan 14. PubMed PMID: 23319141; PubMed Central PMCID: PMC3591667.

## 2012

31. Samel A, Cuomo A, **Bonaldi T**, Ehrenhofer-Murray AE. Methylation of CenH3 arginine 37 regulates kinetochore integrity and chromosome segregation. *Proc Natl Acad Sci U S A.* **2012** Jun 5;109(23):9029-34. doi: 10.1073/pnas.1120968109. Epub 2012 May 21. PubMed PMID: 22615363; PubMed Central PMCID: PMC3384136.

## 2011

32. Vella P, Barozzi I, Cuomo A, **Bonaldi T**, Pasini D. Yin Yang 1 extends the Myc-related transcription factors network in embryonic stem cells. *Nucleic Acids Res.* **2012** Apr;40(8):3403-18. doi: 10.1093/nar/gkr1290. Epub 2011 Dec 30. PubMed PMID: 22210892; PubMed Central PMCID: PMC3333890.
33. Celona B, Weiner A, Di Felice F, Mancuso FM, Cesarini E, Rossi RL, Gregory L, Baban D, Rossetti G, Grianti P, Pagani M, **Bonaldi T**, Ragoussis J, Friedman N, Camilloni G, Bianchi ME, Agresti A. Substantial histone reduction modulates genomewide nucleosomal occupancy and global transcriptional output. *PLoS Biol.* **2011** Jun;9(6):e1001086. doi: 10.1371/journal.pbio.1001086. Epub 2011 Jun 28. PubMed PMID: 21738444; PubMed Central PMCID: PMC3125158.
34. Cuomo A, Moretti S, Minucci S, **Bonaldi T**. SILAC-based proteomic analysis to dissect the "histone modification signature" of human breast cancer cells. *Amino Acids.* **2011** Jul;41(2):387-99. doi: 10.1007/s00726-010-0668-2. Epub 2010 Jul 9. PubMed PMID: 20617350.

## 2009

35. Loyola A, Tagami H, **Bonaldi T**, Roche D, Quivy JP, Imhof A, Nakatani Y, Dent SY, Almouzni G. The HP1 $\alpha$ -CAF1-SetDB1-containing complex provides H3K9me1 for Suv39-mediated K9me3 in pericentric heterochromatin. *EMBO Rep.* **2009** Jul;10(7):769-75. doi: 10.1038/embor.2009.90. Epub 2009 Jun 5. PubMed PMID:19498464; PubMed Central PMCID: PMC2727428.
36. Hilger M, **Bonaldi T**, Gnad F, Mann M. Systems-wide analysis of a phosphatase knock-down by quantitative proteomics and phosphoproteomics. *Mol Cell Proteomics.* **2009** Aug;8(8):1908-20. doi: 10.1074/mcp.M800559-MCP200. Epub 2009 May 9. PubMed PMID: 19429919; PubMed Central PMCID: PMC2722773.

### 2008

37. **Bonaldi T**, Straub T, Cox J, Kumar C, Becker PB, Mann M. Combined use of RNAi and quantitative proteomics to study gene function in Drosophila. *Mol Cell*. **2008** Sep 5;31(5):762-72. doi: 10.1016/j.molcel.2008.07.018. PubMed PMID: 18775334.
38. Bachi A, **Bonaldi T**. Quantitative proteomics as a new piece of the systems biology puzzle. *J Proteomics*. **2008** Aug 21;71(3):357-67. doi: 10.1016/j.jprot.2008.07.001. Epub 2008 Jul 9. Review. PubMed PMID: 18640294.
39. Nielsen ML, Vermeulen M, **Bonaldi T**, Cox J, Moroder L, Mann M. Iodoacetamide-induced artifact mimics ubiquitination in mass spectrometry. *Nat Methods*. **2008** Jun;5(6):459-60. doi: 10.1038/nmeth0608-459. PubMed PMID: 18511913.

### 2007

40. Ferreira R, Eberharter A, **Bonaldi T**, Chioda M, Imhof A, Becker PB. Site-specific acetylation of ISWI by GCN5. *BMC Mol Biol*. **2007** Aug 30;8:73. PubMed PMID: 17760996; PubMed Central PMCID: PMC2045673.

### 2006

41. Loyola A\*, **Bonaldi T\***, Roche D, Imhof A, Almouzni G. PTMs on H3 variants before chromatin assembly potentiate their final epigenetic state. *Mol Cell*. **2006** Oct 20;24(2):309-16. PubMed PMID: 17052464. \**co-first*