Curriculum Vitae

Personal information

First name/ Surname	**Ferdinando Di Cunto**
Address	Molecular Biotechnology Center, University of Torino, Via Nizza 52, 10126, Torino, Italy
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Nationality	Italian
Date of birth	20/12/1969
Gender	Male

Work experience

Dates	1996-1997
Occupation or position held	Research fellow
Main activities and responsibilities	Research activity
Name and address of employer	Massachusetts General Hospital, Boston, Massachusetts, USA
Type of business or sector	Biomedical Research

Dates	1999-2005
Occupation or position held	Researcher (Molecular Biology)
Main activities and responsibilities	Research and teaching
Name and address of employer	University of Torino, Italy
Type of business or sector	Biomedical Research

Dates	2005-2011
Occupation or position held	Associate Professor (Molecular Biology)
Main activities and responsibilities	Research and teaching in Molecular and Computational biology
Name and address of employer	University of Torino, Italy
Type of business or sector	Biomedical Research

Dates	2011-present
Occupation or position held	Full Professor (Molecular Biology)
Main activities and responsibilities	Research and teaching in Molecular and Computational biology

Name and address of employer	University of Torino, Italy
Type of business or sector	Biomedical Research

Education and training

Dates	July 1994
Title of qualification awarded	MD degree (Medicine and Surgery)
Principal subjects/occupational skills covered	Medicine
Name and type of organisation providing education and training	University of Torino

Dates	February 2001
Title of qualification awarded	PhD in “Human Biology, molecular and cellular bases"
Principal subjects/occupational skills covered	Biomedical research
Name and type of organisation providing education and training	University of Torino

Personal skills and competences

Mother tongue(s) **Italian**

Other language(s)

Self-assessment	Understanding		Speaking		Writing
European level (*)	Listening	Reading	Spoken interaction	Spoken production	
English	C2	C2	C2	C2	C2
French	B1	B1	B1	B1	B1

(*) Common European Framework of Reference for Languages

Research interests

1. Study of cytokinesis and of neuronal differentiation during development of the Central Nervous System.
2. Study of the molecular bases of microcephalies
3. Development of computational approaches for the annotation of gene function

Publications

1. Bergo A, Strollo M, Gai M, Barbiero I, Stefanelli G, Sertic S, Cobolli Gigli C, Di Cunto F, Kilstrup-Nielsen C, Landsberger N. (2015). Methyl-CpG binding protein 2 (MeCP2) localizes at the centrosome and is required for proper mitotic spindle organization. J Biol Chem 290:3223-3237.

2. Berto, G. E., Iobbi, C., Camera, P., Scarpa, E., Iampietro, C., Bianchi, F., Gai, M., Sgrò, F., Cristofani, F., Gärtner, A., Dotti, C. G., and Di Cunto, F. (2014). The DCR protein TTC3 affects differentiation and Golgi compactness in neurons through specific actin-regulating pathways. PLoS One 9:e93721.

3. Repetto, D., Camera, P., Melani, R., Morello, N., Russo, I., Calcagno, E., Tomasoni, R., Bianchi, F., Berto, G., Giustetto, M., Berardi, N., Pizzorusso, T., Matteoli, M., Di Stefano, P., Missler, M., Turco, E., Di Cunto, F., and Defilippi, P. (2014). p140Cap regulates memory and synaptic plasticity through Src- and CitN-mediated actin reorganization. Journal of Neuroscience, 34(4):1542-53.

4. Di Gregorio, E., Bianchi, F. T., Schiavi, A., Chiotto, A. M., Rolando, M., Verdun di Cantogno, L., Grosso, E., Cavalieri, S., Calcia, A., Lacerenza, D., Zuffardi, O., Retta, S. F., Stevanin, G., Marelli, C., Durr, A., Forlani, S., Chelly, J., Montarolo, F., Tempia, F., Beggs, H. E., Reed, R., Squadrone, S., Abete, M. C., Brussino, A., Ventura, N., Di Cunto, F., and Brusco, A. (2013). A de novo X;8 translocation creates a PTK2-THOC2 gene fusion with THOC2 expression knockdown in a patient with psychomotor retardation and congenital cerebellar hypoplasia. J Med Genet 50, 543-551.

5. Fagoonee, S., Bearzi, C., Di Cunto, F., Clohessy, J. G., Rizzi, R., Reschke, M., Tolosano, E., Provero, P., Pandolfi, P. P., Silengo, L., and Altruda, F. (2013). The RNA Binding Protein ESRP1 Fine-Tunes the Expression of Pluripotency-Related Factors in Mouse Embryonic Stem Cells. PLoS ONE 8, e72300.

6. Molineris, I., Ala, U., Provero, P., and Di Cunto, F. (2013). Drug repositioning for orphan genetic diseases through Conserved Anticoexpressed Gene Clusters (CAGCs). BMC Bioinformatics 14, 288.

7. Pavan, S., Musiani, D., Torchiaro, E., Migliardi, G., Gai, M., Di Cunto, F., Erriquez, J., Olivero, M., and Di Renzo, M. F. (2013). HSP27 is required for invasion and metastasis triggered by hepatocyte growth factor. Int J Cancer.

8. Piro, R. M., Molineris, I., Di Cunto, F., Eils, R., and Konig, R. (2013). Disease-gene discovery by integration of 3D gene expression and transcription factor binding affinities. Bioinformatics 29, 468-475.

9. Spaccarotella, E., Pellegrino, E., Ferracin, M., Ferreri, C., Cuccuru, G., Liu, C., Iqbal, J., Cantarella, D., Taulli, R., Provero, P., Di Cunto, F., Medico, E., Negrini, M., Chan, W. C., Inghirami, G., and Piva, R. (2013). STAT3-mediated activation of microRNA cluster 17~92 promotes proliferation and survival of ALK positive anaplastic large cell lymphoma. Haematologica.

10. Ugolotti, R., Mesejo, P., Zongaro, S., Bardoni, B., Berto, G., Bianchi, F., Molineris, I., Giacobini, M., Cagnoni, S., and Di Cunto, F. (2013). Visual Search of Neuropil-Enriched RNAs from Brain In Situ Hybridization Data through the Image Analysis Pipeline Hippo-ATESC. PLoS ONE 8, e74481.

11. Piro, R. M., and Di Cunto, F. (2012a). Computational approaches to disease-gene prediction: rationale, classification and successes. Febs J 279, 678-696.

12. Piro, R. M., Molineris, I., Ala, U., and Di Cunto, F. (2012b). Evaluation of candidate genes from orphan FEB and GEFS+ loci by analysis of human brain gene expression atlases. PLoS ONE 6, e23149.

13. Bosio, Y., Berto, G., Camera, P., Bianchi, F., Ambrogio, C., Claus, P., and Di Cunto, F. (2012). PPP4R2 regulates neuronal cell differentiation and survival, functionally cooperating with SMN. Eur J Cell Biol 91, 662-674.

14. Lembo, A., Di Cunto, F., and Provero, P. (2012). Shortening of 3'UTRs correlates with poor prognosis in breast and lung cancer. PLoS ONE 7, e31129.

15. Bosio Y, Berto G, Camera P, Bianchi F, Ambrogio C, Claus P, Di Cunto F. 2012. PPP4R2 regulates neuronal cell differentiation and survival, functionally cooperating with SMN. European Journal of Cell Biology In Press.

16. Piro RM, Di Cunto F. 2012. Computational approaches to disease gene prediction: rationale, classification and successes. Febs J.

17. Tay Y, Kats L, Salmena L, Weiss D, Tan SM, Ala U, Karreth F, Poliseno L, Provero P, Di Cunto F, Lieberman J, Rigoutsos I, Pandolfi PP. 2011. Coding-Independent Regulation of the Tumor Suppressor PTEN by Competing Endogenous mRNAs. Cell 147:344-357.

18. Bianchi FT, Camera P, Ala U, Imperiale D, Migheli A, Boda E, Tempia F, Berto G, Bosio Y, Oddo S, LaFerla FM, Taraglio S, Dotti CG, Di Cunto F. 2011. The collagen chaperone HSP47 is a new interactor of APP that affects the levels of extracellular beta-amyloid peptides. PLoS One 6:e22370.

19. Gai M, Camera P, Dema A, Bianchi F, Berto G, Scarpa E, Germena G, Di Cunto F. 2011. Citron kinase controls abscission through RhoA and Anillin. Mol Biol Cell.

20. Piro R, Molineris I, Ala U, Di Cunto F. 2011. Evaluation of candidate genes from orphan FEB and GEFS+ loci by analysis of human brain gene expression atlases.

21. Piro RM, Ala U, Molineris I, Grassi E, Bracco C, Perego GP, Provero P, Di Cunto F. 2011. An atlas of tissue-specific conserved coexpression for functional annotation and disease gene prediction. Eur J Hum Genet.

22. Damasco, C., Lembo, A., Somma, M. P., Gatti, M., Di Cunto, F., and Provero, P. (2011). A Signature Inferred from Drosophila Mitotic Genes Predicts Survival of Breast Cancer Patients. PLoS ONE 6, e14737.

23. Molineris, I., Grassi, E., Ala, U., Di Cunto, F., and Provero, P. (2011). Evolution of promoter affinity for transcription factors in the human lineage. Mol Biol Evol.

24. Amoresano, A., Di Costanzo, A., Leo, G., Di Cunto, F., La Mantia, G., Guerrini, L., and Calabro, V. (2010). Identification of DeltaNp63alpha protein interactions by mass spectrometry. J Proteome Res 9, 2042-2048.

25. Forlani, G., Giarda, E., Ala, U., Di Cunto, F., Salani, M., Tupler, R., Kilstrup-Nielsen, C., and Landsberger, N. (2010). The MeCP2/YY1 interaction regulates ANT1 expression at 4q35: novel hints for Rett syndrome pathogenesis. Hum Mol Genet 19, 3114-3123.

26. Molineris, I., Sales, G., Bianchi, F., Di Cunto, F., and Caselle, M. (2010). A new approach for the identification of processed pseudogenes. J Comput Biol 17, 755-765.

27. Piro, R. M., Molineris, I., Ala, U., Provero, P., and Di Cunto, F. (2010). Candidate gene prioritization based on spatially mapped gene expression: an application to XLMR. Bioinformatics 26, i618-624.

28. Roetto, A., Di Cunto, F., Pellegrino, R. M., Hirsch, E., Azzolino, O., Bondi, A., Defilippi, I., Carturan, S., Miniscalco, B., Riondato, F., Cilloni, D., Silengo, L., Altruda, F., Camaschella, C., and Saglio, G. (2010). Comparison of 3 Tfr2-deficient murine models suggests distinct functions for Tfr2-alpha and Tfr2-beta isoforms in different tissues. Blood 115, 3382-3389.

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31. Ala, U., Piro, R. M., Grassi, E., Damasco, C., Silengo, L., Oti, M., Provero, P., and Di Cunto, F. (2008). Prediction of human disease genes by human-mouse conserved coexpression analysis. PLoS Comput Biol 4, e1000043.

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33. Camera, P., Schubert, V., Pellegrino, M., Berto, G., Vercelli, A., Muzzi, P., Hirsch, E., Altruda, F., Dotti, C. G., and Di Cunto, F. (2008). The RhoA-associated protein Citron-N controls dendritic spine maintenance by interacting with spine-associated Golgi compartments. EMBO Rep 9, 384-392.

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37. Fagoonee S, Di Cunto F, Vozzi D, Volinia S, Pellegrino M, Gasparini P, Silengo L, Altruda F, Tolosano E. Microarray and large-scale in silico--based identification of genes functionally related to Haptoglobin and/or Hemopexin. DNA Cell Biol. 25, 323-30 ,2006

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39. Cora, D., Herrmann, C., Dieterich, C., Di Cunto, F., Provero, P. &Caselle, M. Ab initio identification of putative human transcription factor binding sites by comparative genomics. BMC Bioinformatics 6, 110, 2005.

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motifs to gene expression: a computational approach to regulatory element discovery
in eukaryotes. BMC Bioinformatics 3(1):72002.

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apoptosis. Neuron 28, 115-127, 2000.

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cycle control. Science 280, 1069-1072, 1998.

51. Di Cunto, F., Calautti, E., Hsiao, J., Ong, L., Topley, G., Turco, E. &Dotto, G. P.
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Rho-Rac-binding protein Citron. J Biol Chem 273, 29706-29711, 1998.

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p21Cip1/WAF1 alters keratinocyte growth and differentiation and promotes ras-tumor
progression. Genes Dev 10, 3065-3075, 1996.

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factors beta 2 and beta 3 down-modulation in 12-O-
tetradecanoylphorbol-13-acetate-induced skin hyperplasia. Cancer Res 53,
5517-5522., 1993

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