



THE EUROPEAN SOCIETY
OF HUMAN GENETICS



EUROPEAN HUMAN GENETICS CONFERENCE 2014

in conjunction with the
European Meeting on Psychosocial Aspects of Genetics 2014

May 31 - June 3, Milan, Italy



PROGRAMME



Time	Gold Room	Space 3+4	Brown 3	Brown 1+2	Space 1
cont.	C18 Large scale genomics Chair: O. Zuffardi, H. Scheffer	C19 Internal organs Chair: M. Zollino, B. Melegh	C20 Basic mechanisms in genetics Chair: B. Franco, S. Lyonnet	C21 Rasopathies and CDG Chair: F. Sangiuolo, K. Writzl	C22 Returning results: Ethical and legal issues, joint with EMPAG Chair: F. Faravelli, M. Cornel
11.45	C18.4 Planar cell polarity gene mutations contribute to the etiology of human Neural Tube Defects <i>Patrizia De Marco, E. Merello, G. Piatelli, A. Cama, Z. Kibar, V. Capra; Genova, Italy</i>	C19.4 Identification and functional characterization of ESR2, a new disease gene for 46,XY disorders of sex development (DSD) <i>Dorien Baetens*, T. Guran, L. De Cauwer, L. Looijenga, K. De Bosscher, M. Cools, E. De Baere; Ghent, Belgium</i>	C20.4 Pseudoautosomal region 1 length polymorphism in the human population <i>Martin A. Mensah*, M.S. Hestand, M.H.D. Lamuseau, M. Isrie, N. Vanderheyden, M. Declercq, E.L. Souche, J. Van Houdt, R. Stoeva, H. Van Esch, K. Devriendt, T. Voet, R. Decorte, P.N. Robinson, J.R. Vermeesch; Leuven, Belgium</i>	C21.4 A mutation in PAK3 with a dual molecular effect deregulates the RAS/MAPK pathway and drives an X-linked syndromic phenotype <i>Pamela Magini*, T. Pippucci, I. Tsai, S. Coppola, E. Stellacci, A. Bartoletti-Stella, D. Turchetti, C. Graziano, G. Cenacchi, I. Neri, D.M. Cordelli, V. Marchiani, R. Bergamaschi, G. Gasparre, G. Neri, L. Mazzanti, A. Patrizi, E. Franzoni, G. Romeo, D. Bordo, M. Tartaglia, N. Katsanis, M. Serj; Bologna, Italy</i>	C22.4 International views on sharing incidental findings from whole genome research <i>Anna Middleton, M. Parker, C. Wright, H. Firth, E. Bragin, M. Hurles, O. DDD Project; Cambridge, United Kingdom</i>
12.00	C18.5 Clinical exome sequencing for cerebellar ataxia and spastic paraplegia reveals novel gene-disease associations and uncovers unanticipated rare disorders <i>Erik-Jan Kamsteeg, B.P. van de Warrenburg, S.T. de Bot, M.A.A.P. Willemsen, S. Vermeer, M.I. Schouten, R. Meijer, M. Pennings, C. Gilissen, H. Scheffer; Nijmegen, Netherlands</i>	C19.5 LRP5 variants associated with development of polycystic kidney and liver disease <i>Wybrich R. Cnossen*, R.H.M. te Morsche, A. Hoischen, C. Gilissen, H. Venselaar, S. Mehdi, C. Bergmann, M. Losekoot, M.H. Breuning, D.J.M. Peters, J.A. Veltman, J.P.H. Drenth; Nijmegen, Netherlands</i>	C20.5 Comparative proteomic analysis of different fragile X syndrome cell lines <i>S. Lanni, F. Palumbo, M. Goracci, G. Mancano, A. Vitali, V. Marzano, F. Iavarone, F. Vincenzoni, M. Castagnola, P. Chiurazzi, Elisabetta Tabolacci, G. Neri; Rome, Italy</i>	C21.5 Activating mutations in RRAS underlie a phenotype within the RASopathy spectrum and contribute to leukaemogenesis <i>Francesca Pantaleoni*, M. Jaiswal, E. Flex, S. Martinelli, M. Strullu, E.K. Fansa, A. Caye, A. De Luca, F. Lepri, L. Pannone, S. Paolacci, G. Bocchinfuso, C. Rossi, A. Farrotti, O. Fenneteau, B. Brethon, P. Cianci, E. Di Schiavi, A. Selicorni, B. Dallapiccola, I.C. Cirstea, L. Stella, M. Zenker, B.D. Gelb, H. Cavé, M.R. Ahmadian, M. Tartaglia; Roma, Italy</i>	C22.5 Newborn screenings and whole genome sequencing: the real need of a genuine public involvement <i>Marta Tomasi, A. Santosuosso; Trento, Italy</i>
12.15	C18.6 WES detects disease causing SNVs and CNVs in Primary immunodeficiencies <i>Hanne S. Sorte, A. Stray-Pedersen, P.S. Samarakoon, L. Forbes, T. Gambin, O.K. Rødningen, I.C. Hanson, L.M. Noroski, C. Davis, F. Seeborg, S.K. Nicholas, J.W. Caldwell, N.Y. Chokshi, D. Bayer, C.R. Beck, T.J. Vece, W. Wiszniewski, S.J. Penney, S.N. Jhangiani, D. Muzny, L.O. Mæhle, A. Patel, H.C. Erichsen, T.G. Abrahamson, J. Buchner, G.E. Tjonnfjord, P. Aukrust, L.T. Osnes, M.A. Kulseth, D.E. Undlien, W.T. Shearer, B. Fevang, R.A. Gibbs, R. Lyle, J.S. Orange, J.R. Lupski; Oslo, Norway</i>	C19.6 Digenic model in Alport syndrome <i>Maria Antonietta Mencarelli*, M. van Geel, H. Storey, C. Fallerini, L. Dosa, M. Antonucci, F. Cetta, A. van den Wijngaard, S. Yau, F. Mari, M. Bruttini, F. Ariani, K. Dahan, B. Smeets, F. Flinter, A. Renieri; Siena, Italy</i>	C20.6 RNA-DNA Differences in Endoplasmic Reticulum Stress Response <i>Allison L. Richards*, S. Liu, Z. Zhu, V.G. Cheung; Ann Arbor, United States</i>	C21.6 A New Mouse Model for Costello Syndrome <i>Tania Sorg, B. Arveiler, M. Birling, G. Bou About, M. Champy, F. Dupuy, I. Goncalves, M. Jagla, H. Jacobs, H. Meziane, G. Pavlovic, N. Philip, F. Radvanyi, R. Rossignol, M. Roux, S. Sigaudy, Y. Herault, D. Lacombe; Illkirch, France</i>	C22.6 Current Developments in the Regulation of Direct-to-Consumer Genetic Testing in Europe <i>Louiza M. Kalokairinou*, H.C. Howard, P. Borry; Leuven, Belgium</i>
12.30	Lunch Break on Level 1 & 2				
13.30					

Presentations highlighted by an asterisk (*) and a grey background are from Young Investigator Award Finalists.

PROGRAMME EMPAG 2014 - TUESDAY

17:30 - 19:00	EES2 - EMPAG Educational Session: Qualitative and quantitative methods in psychosocial research	Room Amber 3+4
Chair:	B. Ignacio, C. Bjorvatn	
	EES2.1 Qualitative and quantitative methods in psychosocial research K. O'Doherty, <i>Guelph, Canada</i>	
	Bettina Meiser; <i>Randwick, Australia</i>	
20:30	Networking party	

Tuesday, June 3, 2014

09:00 - 10:30	ESHG-ASHG Building Bridges Session PL3: „Towards finding global agreement on...“ What IF... (Incidental Findings), an interactive Debate - joint with ESHG	Gold Room
Moderator:	Han Brunner, The Netherlands	
Discussants:	<ul style="list-style-type: none">• Angus Clarke, <i>United Kingdom</i>• Martina Cornel, <i>The Netherlands</i>• Robert Green, <i>United States</i>• Stephen Kingsmore, <i>United States</i>• Marjolijn Kriek, <i>The Netherlands</i>• Arnold Munnich, <i>France</i>	
10:30 - 11:00	Coffee break	
11:00 - 12:30	C22 - Returning results: Ethical and legal issues, joint with ESHG	Space 1
Chair:	F. Faravelli, M. Cornel	
11:00	C22.1 The impact of reporting exome and whole genome sequencing: Predicted frequencies of primary, secondary and incidental findings based on modelling Leslie Burnett, <i>L.C. Ding, R.M. Lew, D. Chesher, A.L. Proos;</i> <i>Sydney, Australia</i>	
11:15	C22.2 Defending the child's right to an open future concerning genetic information. Annelien L. Bredenoord, <i>M.C. de Vries, J.J. van Delden;</i> <i>Utrecht, Netherlands</i>	
11:30	C22.3 Implementation of a duty-to-recontact system in molecular and clinical genetics: perspectives from professionals and patients Mirjam Plantinga, <i>W. Lamers, A.V. Ranchor, M.A. Verkerk, E. Birnie, I.M. van Langen;</i> <i>Groningen, Netherlands</i>	
11:45	C22.4 International views on sharing incidental findings from whole genome research Anna Middleton, <i>M. Parker, C. Wright, H. Firth, E. Bragin, M. Hurler, O. DDD Project;</i> <i>Cambridge, United Kingdom</i>	
12:00	C22.5 Newborn screenings and whole genome sequencing: the real need of a genuine public involvement Marta Tomasi, <i>A. Santosuosso;</i> <i>Trento, Italy</i>	
12:15	C22.6 Current Developments in the Regulation of Direct-to-Consumer Genetic Testing in Europe Louiza M. Kalokairinou, <i>H.C. Howard, P. Borry;</i> <i>Leuven, Belgium</i>	
	End of Meeting	



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Final Programme