

World Journal of *Medical Genetics*

World J Med Genet 2013 August 27; 3(3): 9-13



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2011-2015

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MINIREVIEWS

- 9 Genetic counselling in post-genomic era-to be or not to be
Nenad B, Maurizio M

APPENDIX I-V Instructions to authors

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World Journal of Medical Genetics (World J Med Genet, WJMG, online ISSN 2220-3184, DOI: 10.5496) is a peer-reviewed open access academic journal that aims to guide clinical practice and improve diagnostic and therapeutic skills of clinicians.

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INDEXING/ABSTRACTING *World Journal of Medical Genetics* is now indexed in Digital Object Identifier.

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NAME OF JOURNAL
World Journal of Medical Genetics

ISSN
 ISSN 2220-3184 (online)

LAUNCH DATE
 December 27, 2011

FREQUENCY
 Quarterly

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PUBLISHER
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 Flat C, 23/F, Lucky Plaza,
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 Fax: +852-6555-7188
 Telephone: +852-3177-9906
 E-mail: bpgoffice@wjgnet.com
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PUBLICATION DATE
 August 27, 2013

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 Full instructions are available online at http://www.wjgnet.com/2220-3184/g_info_20100722180909.htm

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Genetic counselling in post-genomic era-to be or not to be

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Received: June 3, 2013 Revised: July 27, 2013

Accepted: August 17, 2013

Published online: August 27, 2013

Abstract

With the surge of genetic tests and technologies, genetic counsellors are faced with the challenge of translating emerging scientific knowledge into practical information for patients, clinicians and public health policy makers. The new tests and technologies also are associated with new psychosocial and ethical considerations. New guidelines are needed for each new discovery of the genomic impact on phenotype, pathology and disease while "old" syndromes and "old" pathology, continue to require attention. In the new post-Human Genome Project era, genetic counsellors will be an integral part of translating genomic discoveries into beneficial impact on human disease, health care, and medical benefits. The needs for genetic counselling should be designed into genomic research at the onset. Genetic counsellors need to handle old while rapidly assimilating new information and the principal challenge is to be up to date and updated.

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Key words: Genetic counseling; Clinical application; Translating emerging scientific knowledge; Direct-to-

consumer genetic testing; Clinicians and public health policy makers

Core tip: This paper aims at discussing the aspects and challenges which have to be faced during genetic counselling in the new post-Human Genome Project era with beneficial impact on human disease, health care, and medical benefits. With the surge of genetic tests and technologies, genetic counsellors are faced with the challenge of translating emerging scientific knowledge into practical information for patients, clinicians same as for public health policy makers and the needs for genetic counselling should be designed into genomic research at the onset. Genetic counsellors need to handle old while rapidly assimilating new information.

Nenad B, Maurizio M. Genetic counselling in post-genomic era-to be or not to be. *World J Med Genet* 2013; 3(3): 9-13 Available from: URL: <http://www.wjgnet.com/2220-3184/full/v3/i3/9.htm> DOI: <http://dx.doi.org/10.5496/wjmg.v3.i3.9>

INFLUENCE ON GENETIC COUNSELLING/ COUNSELLORS AND OTHER HEALTH PROFESSIONALS

The spectacular progress in understanding the genetic nature of disease has deeply changed the daily practice of medicine. With the surge of genetic tests and technologies, genetic counsellors are faced with the challenge of translating emerging scientific knowledge into practical information for patients, clinicians and public health policy makers. The new tests and technologies also are associated with new psychosocial and ethical considerations. As reported by Bennett *et al*^[1], the field of genetic counselling arose from the need to educate, manage and counsel individuals and their families diagnosed with, or at risk for, genetic diseases with respect to how these conditions may affect the psychological, medical, finan-

cial and social aspects of one's life.

Genetic counselling is now a necessary component of the practice of virtually all medical specialties. Physicians must help their patients understand a genetic diagnosis and assist them in making and coping with decisions relating to the diagnosis. As each new genetic test is made available in the clinic, developing the appropriate counselling for each new diagnosis is necessarily a multidisciplinary endeavour that includes involvement of a Specialist of Medical Genetics^[2,3]. This paper aims at discussing the aspects and challenges which have to be faced during genetic counselling.

WHAT'S NEW?

The Human Genome Project provide us not only with information regarding the basic architecture of human genome, it also gave rise to impressive advances in molecular technologies. It is now possible to routinely assess genetic variation at a population level. For example, it is routine to assess over a million single nucleotide polymorphisms (SNPs) on thousands of individuals within a single study and it is routine to combine studies into meta-analyses across hundreds of thousands of individuals^[4]. An excellent review of Gao *et al*^[4] discussed the use of genome-wide association studies (GWAS). This strategy is based upon the common disease common variant hypothesis^[5], in which it is proposed that high-prevalence traits are determined by high-frequency genetic variants. Some successful examples are given by meta-analyses in GWAS in Parkinson's disease^[6], type 2 diabetes^[7,8], type 1 diabetes^[9], chronic kidney disease^[10], retinal microcirculation^[11], Crohn's disease^[12], and others. Beyond simply examining nucleotide variations, new technology allows researchers to assess other aspects of genomic variations including whole transcriptome profiling and genome-wide epigenetic modifications. Now the major challenge in genomics is to apply this rapidly expanding plethora of genetic data in meaningful ways-to further improve our understanding of human biology^[13] and to generate knowledge about the genetic contribution to human diseases^[14].

While research focuses on how to put the human genome in context, it should not be forgotten that it is quite "tricky" to translate these research data into appropriate genetic counselling of each client. Especially if we consider that the next step in personal genomics is to associate an individual's specific variation with clinical disease phenotypes, counselling must help individuals discriminate between medically important variation and benign polymorphic variation. Data of genomic variations must be carefully translated by a genetic counsellor with care to educate the clients of the presumed significance of genes and mutations, imprinting, and the likelihood of benign versus causative genomic changes^[15,16]. This means that genetic counsellors are at the forefront of introducing and applying the advances from genome

science to the lives of individuals and their families, by translating the complex language of genomic medicine into terms that are easy to understand^[1].

WHAT HAS BEEN CHANGED AS CONSEQUENCE OF NEW DIAGNOSTIC APPROACHES?

The era of genomic medicine challenges traditional definitions of "healthy" and "diseased". Traditionally, medical attention is only sought regarding a present illness. Now genetic testing permits the diagnosis of healthy individuals who are expected to develop or have an increased susceptibility to develop a disorder^[17,18]. Testing for susceptibility genes will push into the world of medicine millions of individuals who have no personal experience of any disease, as emphasized by Professor Dallapiccola^[19]. Some of them will benefit from the information, but many will become "unpatients", *i.e.*, individuals who are neither patients under treatment nor healthy individuals free of any medically relevant condition. This new class of individuals (it seems appropriate to call them "clients") who are watching and waiting for a sign of disease must be advised to undertake appropriate systematic clinical and instrumental monitoring while avoiding the development of psychosomatic symptoms. It is thus necessary to rethink the genetics revolution in medicine in terms of benefits and harm considering that the general rule for all physicians is: "*Primum non nocere*" ("First, do no harm"). After all, when we think about applications of genetics in daily practice, genetic counselling included, we should take in mind, J. Watson's observation: "I have benefited a lot from being the first human to have his or her personal genome made publicly available on the web. So far, knowledge of my personal genetic risks has not cost me an hour of sleep. I doubt, however, whether I would feel so positively if this knowledge had been given to me at a much earlier stage of my life"^[16,20].

It is necessary to introduce into genomic research consideration of computational strategies which permit translation of genetic information into clinically useful probability estimation. Personalized cancer risk assessment is an example of this integration. Algorithms in conjunction with testing have been successfully applied to predict the probability to carrier germinal at-risk mutations, as BRCAPRO for breast and ovarian cancer syndromes^[21-24], PancPRO for familial pancreatic cancer^[25] and Melapro for melanoma families^[26], *etc.*

ITALIAN EXPERIENCE

The Italian genetic testing survey 2004^[27], could be seen as starting point (at least for us who are working in Italy) for understanding the necessity to link testing and genetic counselling in order to cut the costs, and to widen the number of available services. This survey also stressed the necessity for constant training of the general practi-

tioner and education of the consumer with regard to the appropriate use of genetic tests. A more sparing use of genetic tests, which should always follow specific clinical indications, ideally flank and sustain good clinical practice. Conversely, inappropriate genetics testing do harm by imparting a false sense of reassurance in individuals found not to have a gene mutation who are not informed of the limitation of tests and are major contributors to increasing health care costs^[28,29].

“DO-IT-YOURSELF” GENOMIC TESTING

Direct-to-consumer genetic testing (DTC-GT) provides personalized genetic risk information directly to consumers. DTC-GT has generated a considerable controversy about its potential benefit, harms, and regulatory status since its entry into the mainstream marketplace in 2006^[30,31] largely as a result of the unclear link between DTC-GT results, consumer risk and cost effective health care decisions.

With DTC-GT, clients without the guidance of genetic counselling will often purchase a genetic test that is not clinically indicated. Individuals ordering and interpreting genetic tests for tens or hundreds of conditions with varying clinical validity and utility, in the absence of a healthcare professional, could lead to unnecessary or incorrect healthcare decisions or emotional distress in the clients^[32]. Furthermore afterwards clients may communicate the results to health-care professionals-it is left to these professionals to discuss the testing guidelines and clinical/diagnostic protocols and the usefulness and significance of the results, opening the door to distrust and misunderstandings if the test results are discounted^[33].

Obviously access to DTC-GT can be seen as a right for consumers to purchase the offered product and services. However, the open issues about whether and how to regulate the new heterogeneous DTC-GT industry should be systematically and carefully studied to ascertain the clinical utility, referral patterns and downstream costs^[32,33].

SOME PERSPECTIVES WHICH ARE BECOMING REALITY, IS THIS SCIENCE OR SCIENCE FICTION?

Autism spectrum disorders (ASDs) are an example of an emerging area for genomic diagnosis that will require parallel development of genetic counselling. ADS are a heterogeneous group of neurodevelopmental disorders affecting social communication, language and behavior.

There have been reports of applying panels of common SNPs to assess ASD risk^[34,35], but these approaches require more testing/investigation before SNPs can be associated with risk. With rapid emergence of whole-genome sequencing studies, there will be an explosion of new data leading to more comprehensive genotype and phenotype studies^[36-38]. In addition to seeking to identify

genes that influence diseases, scientists are looking for genes which influence biological markers of disease or endophenotype. One example of this approach is the emerging field of imaging genomics which discover important variants associated with brain structure and function. In these studies, a high degree of correlation has been observed between genome and image-derived maps giving some explanation on how these variations influence disease risk and fundamental cognitive processes^[39].

Parents of children with ASDs are generally aware that their subsequent children are at increased risk to be ‘on the spectrum’, but parents often over- or underestimate this risk. While the studies to date indicate that it may be possible, as yet no definitive genomic diagnostic or prognostic indicators of ASD have been found that can be used for risk estimation. The genetic testing and counselling approach to individuals with ASDs will continue to evolve as we learn more about the genetic factors involved and their relative contributions. The next step is to interpret this data and translate it in comprehensive and useful genetic counselling.

LAST BUT NOT LEAST

The scope of genomic counselling are growing rapidly. New guidelines are needed for each new discovery of the genomic impact on phenotype, pathology and disease while “old” syndromes and “old” pathology, for example Downs Syndrome, continue to require attention. That is one of the reasons why the guidelines such as those published by National Society of Genetic Counsellors-Sheets *et al*^[40], will be always welcomed and “evergreen”.

In the new post-Human Genome Project era, genetic counsellors will be an integral part of translating genomic discoveries into beneficial impact on human disease, health care, and medical benefits. The needs for genetic counselling should be designed into genomic research at the onset. Genetic counsellors need to handle old while rapidly assimilating new information. The principal challenge is to be up to date and updated.

ACKNOWLEDGMENTS

The authors would like to thank John Elling, PhD, for his useful and constructive comments and suggestions on the manuscript.

REFERENCES

- 1 **Bennett RL**, Hampel HL, Mandell JB, Marks JH. Genetic counselors: translating genomic science into clinical practice. *J Clin Invest* 2003; **112**: 1274-1279 [PMID: 14597750 DOI: 10.1172/JCI200320113]
- 2 **Lee FH**, Raja SN. Should anesthesiologists be equipped as genetic counselors? *Anesthesiology* 2010; **113**: 507-509 [PMID: 20683249 DOI: 10.1097/ALN.0b013e3181e89ace]
- 3 **Rosenberg H**, Vladutiu GD, Larach MG. Anesthesiologists as genetic counselors? *Anesthesiology* 2011; **114**: 1003;

- author reply 1003-1004 [PMID: 21427547 DOI: 10.1097/ALN.0b013e31820ef9e2]
- 4 **Gao X**, Edwards TL. Genome-wide association studies: Where we are heading? *World J Med Genet* 2011; 1: 23-35 [DOI: 10.5496/wjmg.v1.i1.23]
 - 5 **International HapMap Consortium**. The International HapMap Project. *Nature* 2003; **426**: 789-796 [PMID: 14685227 DOI: 10.1038/nature02168]
 - 6 **Evangelou E**, Maraganore DM, Ioannidis JP. Meta-analysis in genome-wide association datasets: strategies and application in Parkinson disease. *PLoS One* 2007; **2**: e196 [PMID: 17332845 DOI: 10.1371/journal.pone.0000196]
 - 7 **Scott LJ**, Mohlke KL, Bonnycastle LL, Willer CJ, Li Y, Duren WL, Erdos MR, Stringham HM, Chines PS, Jackson AU, Prokunina-Olsson L, Ding CJ, Swift AJ, Narisu N, Hu T, Pruim R, Xiao R, Li XY, Conneely KN, Riebow NL, Sprau AG, Tong M, White PP, Hetrick KN, Barnhart MW, Bark CW, Goldstein JL, Watkins L, Xiang F, Saramies J, Buchanan TA, Watanabe RM, Valle TT, Kinnunen L, Abecasis GR, Pugh EW, Doheny KF, Bergman RN, Tuomilehto J, Collins FS, Boehnke M. A genome-wide association study of type 2 diabetes in Finns detects multiple susceptibility variants. *Science* 2007; **316**: 1341-1345 [PMID: 17463248 DOI: 10.1126/science.1142382]
 - 8 **Zeggini E**, Scott LJ, Saxena R, Voight BF, Marchini JL, Hu T, de Bakker PI, Abecasis GR, Almgren P, Andersen G, Ardlie K, Boström KB, Bergman RN, Bonnycastle LL, Borch-Johnsen K, Burt NP, Chen H, Chines PS, Daly MJ, Deodhar P, Ding CJ, Doney AS, Duren WL, Elliott KS, Erdos MR, Frayling TM, Freathy RM, Gianniny L, Grallert H, Grarup N, Groves CJ, Guiducci C, Hansen T, Herder C, Hitman GA, Hughes TE, Isomaa B, Jackson AU, Jørgensen T, Kong A, Kubalanza K, Kuruvilla FG, Kuusisto J, Langenberg C, Lango H, Lauritzen T, Li Y, Lindgren CM, Lyssenko V, Marville AF, Meisinger C, Midthjell K, Mohlke KL, Morken MA, Morris AD, Narisu N, Nilsson P, Owen KR, Palmer CN, Payne F, Perry JR, Pettersen E, Platou C, Prokopenko I, Qi L, Qin L, Rayner NW, Rees M, Roix JJ, Sandbaek A, Shields B, Sjögren M, Steinthorsdottir V, Stringham HM, Swift AJ, Thorleifsson G, Thorsteinsdottir U, Timpson NJ, Tuomi T, Tuomilehto J, Walker M, Watanabe RM, Weedon MN, Willer CJ, Illig T, Hveem K, Hu FB, Laakso M, Stefansson K, Pedersen O, Wareham NJ, Barroso I, Hattersley AT, Collins FS, Groop L, McCarthy MI, Boehnke M, Altshuler D. Meta-analysis of genome-wide association data and large-scale replication identifies additional susceptibility loci for type 2 diabetes. *Nat Genet* 2008; **40**: 638-645 [PMID: 18372903 DOI: 10.1038/ng.120]
 - 9 **Barrett JC**, Clayton DG, Concannon P, Akolkar B, Cooper JD, Erlich HA, Julier C, Morahan G, Nerup J, Nierras C, Plagnol V, Pociot F, Schuilenburg H, Smyth DJ, Stevens H, Todd JA, Walker NM, Rich SS. Genome-wide association study and meta-analysis find that over 40 loci affect risk of type 1 diabetes. *Nat Genet* 2009; **41**: 703-707 [PMID: 19430480 DOI: 10.1038/ng.381]
 - 10 **Köttgen A**, Pattaro C, Böger CA, Fuchsberger C, Olden M, Glazer NL, Parsa A, Gao X, Yang Q, Smith AV, O'Connell JR, Li M, Schmidt H, Tanaka T, Isaacs A, Ketkar S, Hwang SJ, Johnson AD, Dehghan A, Teumer A, Paré G, Atkinson EJ, Zeller T, Lohman K, Cornelis MC, Probst-Hensch NM, Kronenberger F, Tönjes A, Hayward C, Aspelund T, Eiriksdottir G, Launer LJ, Harris TB, Rampersaud E, Mitchell BD, Arking DE, Boerwinkle E, Struchalin M, Cavalieri M, Singleton A, Giallauria F, Metter J, de Boer IH, Haritunians T, Lumley T, Siscovick D, Psaty BM, Zillikens MC, Oostra BA, Feitosa M, Province M, de Andrade M, Turner ST, Schillert A, Ziegler A, Wild PS, Schnabel RB, Wilde S, Munzel TF, Leak TS, Illig T, Klopp N, Meisinger C, Wichmann HE, Koenig W, Zgaga L, Zemunik T, Kolcic I, Minelli C, Hu FB, Johansson A, Igl W, Zaboli G, Wild SH, Wright AF, Campbell H, Ellinghaus D, Schreiber S, Aulchenko YS, Felix JF, Rivadeneira F, Uitterlinden AG, Hofman A, Imboden M, Nitsch D, Brandstätter A, Kollerits B, Kedenko L, Mägi R, Stumvoll M, Kovacs P, Boban M, Campbell S, Endlich K, Völzke H, Kroemer HK, Nauck M, Völker U, Polasek O, Vitart V, Badola S, Parker AN, Ridker PM, Kardia SL, Blankenberg S, Liu Y, Curhan GC, Franke A, Roach T, Paulweber B, Prokopenko I, Wang W, Gudnason V, Shuldiner AR, Coresh J, Schmidt R, Ferrucci L, Shlipak MG, van Duijn CM, Borecki I, Krämer BK, Rudan I, Gyllensten U, Wilson JF, Witteman JC, Pramstaller PP, Rettig R, Hastie N, Chasman DI, Kao WH, Heid IM, Fox CS. New loci associated with kidney function and chronic kidney disease. *Nat Genet* 2010; **42**: 376-384 [PMID: 20383146 DOI: 10.1038/ng.568]
 - 11 **Ikram MK**, Sim X, Jensen RA, Cotch MF, Hewitt AW, Ikram MA, Wang JJ, Klein R, Klein BE, Breteler MM, Cheung N, Liew G, Mitchell P, Uitterlinden AG, Rivadeneira F, Hofman A, de Jong PT, van Duijn CM, Kao L, Cheng CY, Smith AV, Glazer NL, Lumley T, McKnight B, Psaty BM, Jonasson F, Eiriksdottir G, Aspelund T, Harris TB, Launer LJ, Taylor KD, Li X, Iyengar SK, Xi Q, Sivakumaran TA, Mackey DA, Macgregor S, Martin NG, Young TL, Bis JC, Wiggins KL, Heckbert SR, Hammond CJ, Andrew T, Fahy S, Attia J, Holliday EG, Scott RJ, Islam FM, Rotter JI, McAuley AK, Boerwinkle E, Tai ES, Gudnason V, Siscovick DS, Vingerling JR, Wong TY. Four novel Loci (19q13, 6q24, 12q24, and 5q14) influence the microcirculation *in vivo*. *PLoS Genet* 2010; **6**: e1001184 [PMID: 21060863 DOI: 10.1371/journal.pgen.1001184]
 - 12 **Franke A**, McGovern DP, Barrett JC, Wang K, Radford-Smith GL, Ahmad T, Lees CW, Balschun T, Lee J, Roberts R, Anderson CA, Bis JC, Bumpstead S, Ellinghaus D, Festen EM, Georges M, Green T, Haritunians T, Jostins L, Latiano A, Mathew CG, Montgomery GW, Prescott NJ, Raychaudhuri S, Rotter JI, Schumm P, Sharma Y, Simms LA, Taylor KD, Whiteman D, Wijmenga C, Baldassano RN, Barclay M, Bayless TM, Brand S, Büning C, Cohen A, Colombel JF, Cottone M, Stronati L, Denson T, De Vos M, D'Inca R, Dubinsky M, Edwards C, Florin T, Franchimont D, Geary R, Glas J, Van Gossom A, Guthery SL, Halfvarson J, Verspaget HW, Hugot JP, Karban A, Laukens D, Lawrance I, Lemann M, Levine A, Libioulle C, Louis E, Mowat C, Newman W, Panés J, Phillips A, Proctor DD, Regueiro M, Russell R, Rutgeerts P, Sanderson J, Sans M, Seibold F, Steinhardt AH, Stokkers PC, Torkvist L, Kullak-Ublick G, Wilson D, Walters T, Targan SR, Brant SR, Rioux JD, D'Amato M, Weersma RK, Kugathasan S, Griffiths AM, Mansfield JC, Vermeire S, Duerr RH, Silverberg MS, Satsangi J, Schreiber S, Cho JH, Annesse V, Hakonarson H, Daly MJ, Parkes M. Genome-wide meta-analysis increases to 71 the number of confirmed Crohn's disease susceptibility loci. *Nat Genet* 2010; **42**: 1118-1125 [PMID: 21102463 DOI: 10.1038/ng.717]
 - 13 **Comuzzie AG**. A grand challenge for applied genetic epidemiology: putting the human genome in context. *Front Genet* 2011; **2**: 10 [PMID: 22303309 DOI: 10.3389/fgene.2011.00010]
 - 14 **van Bokhoven H**. What is the purpose of launching the World Journal of Medical Genetics. *World J Med Genet* 2011; **1**: 1-3 [DOI: 10.5496/wjmg.v1.i1.1]
 - 15 **Wang T**, Pradhan K, Ye K, Wong LJ, Rohan TE. Estimating allele frequency from next-generation sequencing of pooled mitochondrial DNA samples. *Front Genet* 2011; **2**: 51 [PMID: 22303347 DOI: 10.3389/fgene.2011.00051]
 - 16 **Auffray C**, Caulfield T, Khoury MJ, Lupski JR, Schwab M, Veenstra T. Genome Medicine: past, present and future. *Genome Med* 2011; **3**: 6 [PMID: 21345269 DOI: 10.1186/gm220]
 - 17 **Jonsen AR**, Durfy SJ, Burke W, Motulsky AG. The advent of the "unpatients". *Nat Med* 1996; **2**: 622-624 [PMID: 8640544]
 - 18 **Wheeler DA**, Srinivasan M, Egholm M, Shen Y, Chen L, McGuire A, He W, Chen YJ, Makhijani V, Roth GT, Gomes X, Tartaro K, Niazi F, Turcotte CL, Irzyk GP, Lupski JR, Chinault C, Song XZ, Liu Y, Yuan Y, Nazareth L, Qin X, Muzny DM, Margulies M, Weinstock GM, Gibbs RA, Roth-

- berg JM. The complete genome of an individual by massively parallel DNA sequencing. *Nature* 2008; **452**: 872-876 [PMID: 18421352 DOI: 10.1038/nature06884]
- 19 **Bruno Dallapiccola**. The Forum, Genetics and the Future of Europe, Session 4-Responsible Use of Genetics, Challenges. Available from: URL: <http://ec.europa.eu/research/quality-of-life/genetics/en/proceedings-os.html>
- 20 **Watson J**. Living with my personal genome. *Personalized Medicine* 2009; **6**: 607 [DOI: 10.2217/pme.09.62]
- 21 **Parmigiani G**, Berry D, Aguilar O. Determining carrier probabilities for breast cancer-susceptibility genes BRCA1 and BRCA2. *Am J Hum Genet* 1998; **62**: 145-158 [PMID: 9443863 DOI: 10.1086/301670]
- 22 **Berry DA**, Parmigiani G, Sanchez J, Schildkraut J, Winer E. Probability of carrying a mutation of breast-ovarian cancer gene BRCA1 based on family history. *J Natl Cancer Inst* 1997; **89**: 227-238 [PMID: 9017003 DOI: 10.1093/jnci/89.3.227]
- 23 **Antoniou AC**, Gayther SA, Stratton JF, Ponder BA, Easton DF. Risk models for familial ovarian and breast cancer. *Genet Epidemiol* 2000; **18**: 173-190 [PMID: 10642429 DOI: 10.1002/(SICI)1098-2272(200002)18:2<173::AID-GEPI6>3.0.CO;2-R]
- 24 **Nanda R**, Schumm LP, Cummings S, Fackenthal JD, Sveen L, Ademuyiwa F, Cobleigh M, Esserman L, Lindor NM, Neuhausen SL, Olopade OI. Genetic testing in an ethnically diverse cohort of high-risk women: a comparative analysis of BRCA1 and BRCA2 mutations in American families of European and African ancestry. *JAMA* 2005; **294**: 1925-1933 [PMID: 16234499 DOI: 10.1001/jama.294.15.1925]
- 25 **Wang W**, Chen S, Brune KA, Hruban RH, Parmigiani G, Klein AP. PancPRO: risk assessment for individuals with a family history of pancreatic cancer. *J Clin Oncol* 2007; **25**: 1417-1422 [PMID: 17416862 DOI: 10.1200/JCO.2006.09.2452]
- 26 **Wang W**, Niendorf KB, Patel D, Blackford A, Marroni F, Sober AJ, Parmigiani G, Tsao H. Estimating CDKN2A carrier probability and personalizing cancer risk assessments in hereditary melanoma using MelaPRO. *Cancer Res* 2010; **70**: 552-559 [PMID: 20068151 DOI: 10.1158/0008-5472.CAN-09-2653]
- 27 **Dallapiccola B**, Torrente I, Morena A, Dagna-Bricarelli F, Mingarelli R. Genetic testing in Italy, year 2004. *Eur J Hum Genet* 2006; **14**: 911-916 [PMID: 16724000 DOI: 10.1038/sj.ejhg.5201653]
- 28 **Pearson H**. Genetic test adverts under scrutiny. Nature Science Update 2003. Available from: URL: <http://www.nature.com/nsu/030317/030317-3.html>
- 29 **Moreno J**. Selling genetic tests: shades of gray in your DNA. 23 Sept, 2003. Available from: URL: http://abcnews.go.com/sections/living/DailyNews/ONCALL_DTC_brca_tests020923.html
- 30 **Frueh FW**, Greely HT, Green RC, Hogarth S, Siegel S. The future of direct-to-consumer clinical genetic tests. *Nat Rev Genet* 2011; **12**: 511-515 [PMID: 21629275 DOI: 10.1038/nrg3026]
- 31 **Kolor K**, Duquette D, Zlot A, Foland J, Anderson B, Giles R, Wrathall J, Khoury MJ. Public awareness and use of direct-to-consumer personal genomic tests from four state population-based surveys, and implications for clinical and public health practice. *Genet Med* 2012; **14**: 860-867 [PMID: 22814860 DOI: 10.1038/gim.2012.67]
- 32 **Bollinger JM**, Green RC, Kaufman D. Attitudes about regulation among direct-to-consumer genetic testing customers. *Genet Test Mol Biomarkers* 2013; **17**: 424-428 [PMID: 23560882 DOI: 10.1089/gtmb.2012.0453]
- 33 **Giovanni MA**, Fickie MR, Lehmann LS, Green RC, Meckley LM, Veenstra D, Murray MF. Health-care referrals from direct-to-consumer genetic testing. *Genet Test Mol Biomarkers* 2010; **14**: 817-819 [PMID: 20979566 DOI: 10.1089/gtmb.2010.0051]
- 34 **Devlin B**, Scherer SW. Genetic architecture in autism spectrum disorder. *Curr Opin Genet Dev* 2012; **22**: 229-237 [PMID: 22463983 DOI: 10.1016/j.gde.2012.03.002]
- 35 **Skafidas E**, Testa R, Zantomio D, Chana G, Everall IP, Pantelis C. Predicting the diagnosis of autism spectrum disorder using gene pathway analysis. *Mol Psychiatry* 2012 Sep 11; Epub ahead of print [PMID: 22965006 DOI: 10.1038/mp.2012.126]
- 36 **Mercer L**, Creighton S, Holden JJ, Lewis ME. Parental perspectives on the causes of an autism spectrum disorder in their children. *J Genet Couns* 2006; **15**: 41-50 [PMID: 16547798 DOI: 10.1007/s10897-005-9002-7]
- 37 **Selkirk CG**, McCarthy Veach P, Lian F, Schimmenti L, LeRoy BS. Parents' perceptions of autism spectrum disorder etiology and recurrence risk and effects of their perceptions on family planning: Recommendations for genetic counselors. *J Genet Couns* 2009; **18**: 507-519 [PMID: 19488842 DOI: 10.1007/s10897-009-9233-0]
- 38 **Carter MT**, Scherer SW. Autism spectrum disorder in the genetics clinic: a review. *Clin Genet* 2013; **83**: 399-407 [PMID: 23425232 DOI: 10.1111/cge.12101]
- 39 **Hibar DP**, Kohannim O, Stein JL, Chiang MC, Thompson PM. Multilocus genetic analysis of brain images. *Front Genet* 2011; **2**: 73 [PMID: 22303368 DOI: 10.3389/fgene.2011.00073]
- 40 **Sheets KB**, Crissman BG, Feist CD, Sell SL, Johnson LR, Donahue KC, Masser-Frye D, Brookshire GS, Carre AM, Lagrave D, Brasington CK. Practice guidelines for communicating a prenatal or postnatal diagnosis of Down syndrome: recommendations of the national society of genetic counselors. *J Genet Couns* 2011; **20**: 432-441 [PMID: 21618060 DOI: 10.1007/s10897-011-9375-8]

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GENERAL INFORMATION

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WJMG covers topics concerning genes and the pathology of human disease, molecular analysis of simple and complex genetic traits, cancer genetics, epigenetics, gene therapy, developmental genetics, regulation of gene expression, strategies and technologies for extracting function from genomic data, pharmacological genomics, genome evolution. The current columns of *WJMG* include editorial, frontier, diagnostic advances, therapeutics advances, field of vision, mini-reviews, review, topic highlight, medical ethics, original articles, case report, clinical case conference (Clinicopathological conference), and autobiography.

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Name of journal

World Journal of Medical Genetics

ISSN

ISSN 2220-3184 (online)

Launch date

December 27, 2011

Frequency

Quarterly

Editor-in-Chief

Hans van Bokhoven, Professor, PhD, Department of Human

Instructions to authors

Genetics and Cognitive Neurosciences, Radboud university Nijmegen
Medical centre, PO Box 9101, 6500 HB Nijmegen, The Netherlands

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No. 62 Dongsihuan Zhonglu, Chaoyang District,
Beijing 100025, China
Telephone: +86-10-85381891
Fax: +86-10-85381893
E-mail: wjmg@wjgnet.com
<http://www.wjgnet.com>

Publisher

Baishideng Publishing Group Co., Limited
Flat C, 23/F, Lucky Plaza, 315-321 Lockhart Road,
Wan Chai, Hong Kong, China
Telephone: +852-58042046
Fax: +852-31158812
E-mail: bpgoffice@wjgnet.com
<http://www.wjgnet.com>

Production center

Beijing Baishideng BioMed Scientific Co., Limited
Room 903, Building D, Ocean International Center,
No. 62 Dongsihuan Zhonglu, Chaoyang District,
Beijing 100025, China
Telephone: +86-10-85381892
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Instructions to authors

Full instructions are available online at http://www.wjgnet.com/2220-3184/g_info_20100722180909.htm.

Indexed and Abstracted in

Digital Object Identifier.

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Statistical review is performed after peer review. We invite an expert in Biomedical Statistics to evaluate the statistical method used in the paper, including *t*-test (group or paired comparisons), chi-squared test, Ridit, probit, logit, regression (linear, curvilinear, or stepwise), correlation, analysis of variance, analysis of covariance, *etc.* The reviewing points include: (1) Statistical methods should be described when they are used to verify the results; (2) Whether the statistical techniques are suitable or correct; (3) Only homogeneous data can be averaged. Standard deviations are preferred to standard errors. Give the number of observations and subjects (*n*). Losses in observations, such as drop-outs from the study should be reported; (4) Values such as ED50, LD50, IC50 should have their 95% confidence limits calculated and compared by weighted probit analysis (Bliss and Finney); and (5) The word "significantly" should be replaced by its synonyms (if it indicates extent) or the *P* value (if it indicates statistical significance).

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In the interests of transparency and to help reviewers assess any potential bias, *WJMG* requires authors of all papers to declare any competing commercial, personal, political, intellectual, or religious interests

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In press

- 3 **Tian D**, Araki H, Stahl E, Bergelson J, Kreitman M. Signature

of balancing selection in Arabidopsis. *Proc Natl Acad Sci USA* 2006; In press

Organization as author

- 4 **Diabetes Prevention Program Research Group**. Hypertension, insulin, and proinsulin in participants with impaired glucose tolerance. *Hypertension* 2002; **40**: 679-686 [PMID: 12411462 PMID:2516377 DOI:10.1161/01.HYP.0000035706.28494.09]

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- 5 **Vallancien G**, Emberton M, Harving N, van Moorselaar RJ; Alf-One Study Group. Sexual dysfunction in 1, 274 European men suffering from lower urinary tract symptoms. *J Urol* 2003; **169**: 2257-2261 [PMID: 12771764 DOI:10.1097/01.ju.0000067940.76090.73]

No author given

- 6 21st century heart solution may have a sting in the tail. *BMJ* 2002; **325**: 184 [PMID: 12142303 DOI:10.1136/bmj.325.7357.184]

Volume with supplement

- 7 **Geraud G**, Spierings EL, Keywood C. Tolerability and safety of frovatriptan with short- and long-term use for treatment of migraine and in comparison with sumatriptan. *Headache* 2002; **42** Suppl 2: S93-99 [PMID: 12028325 DOI:10.1046/j.1526-4610.42.s2.7.x]

Issue with no volume

- 8 **Banitt DM**, Kaufer H, Hartford JM. Intraoperative frozen section analysis in revision total joint arthroplasty. *Clin Orthop Relat Res* 2002; (**401**): 230-238 [PMID: 12151900 DOI:10.1097/00003086-200208000-00026]

No volume or issue

- 9 Outreach: Bringing HIV-positive individuals into care. *HRS-A Careaction* 2002; 1-6 [PMID: 12154804]

Books

Personal author(s)

- 10 **Sherlock S**, Dooley J. Diseases of the liver and biliary system. 9th ed. Oxford: Blackwell Sci Pub, 1993: 258-296

Chapter in a book (list all authors)

- 11 **Lam SK**. Academic investigator's perspectives of medical treatment for peptic ulcer. In: Swabb EA, Azabo S. Ulcer disease: investigation and basis for therapy. New York: Marcel Dekker, 1991: 431-450

Author(s) and editor(s)

- 12 **Breedlove GK**, Schorfheide AM. Adolescent pregnancy. 2nd ed. Wiczorek RR, editor. White Plains (NY): March of Dimes Education Services, 2001: 20-34

Conference proceedings

- 13 **Harnden P**, Joffe JK, Jones WG, editors. Germ cell tumours V. Proceedings of the 5th Germ cell tumours Conference; 2001 Sep 13-15; Leeds, UK. New York: Springer, 2002: 30-56

Conference paper

- 14 **Christensen S**, Oppacher F. An analysis of Koza's computational effort statistic for genetic programming. In: Foster JA, Lutton E, Miller J, Ryan C, Tettamanzi AG, editors. Genetic programming. EuroGP 2002: Proceedings of the 5th European Conference on Genetic Programming; 2002 Apr 3-5; Kinsdale, Ireland. Berlin: Springer, 2002: 182-191

Electronic journal (list all authors)

- 15 Morse SS. Factors in the emergence of infectious diseases. *Emerg Infect Dis* serial online, 1995-01-03, cited 1996-06-05; 1(1): 24 screens. Available from: URL: <http://www.cdc.gov/ncidod/eid/index.htm>

Patent (list all authors)

- 16 **Pagedas AC**, inventor; Ancel Surgical R&D Inc., assignee. Flexible endoscopic grasping and cutting device and positioning tool assembly. United States patent US 20020103498. 2002 Aug 1

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