Genomic medicine in Africa: promise, problems and prospects

Ambroise Wonkam1,2* and Bongani M Mayosi2

The promise

Remarkable progress has been made in using genomic information to determine how genes are regulated, and how they interact with each other and with the environment to control complex biochemical functions of living organisms in health and disease [1]. This information will have major benefits for the prevention, diagnosis and management of many diseases, including communicable and genetic diseases. In Africa, where infectious diseases are highly prevalent, research on pathogen genomes has enhanced our understanding of disease transmission, virulence mechanisms and avoidance of host defenses [2]. It is anticipated that this information will enable the development of new diagnostic tests, vaccines and therapeutic agents; it is also likely to lead to new approaches for vector control, and reveal why individuals and populations vary in their susceptibility to infectious diseases [1].

This commentary focuses on problems and prospects for the use of new genomic knowledge in improving the health of Africans. The prospects for genomic medicine in Africa have been enhanced by major initiatives that are led by international funding agencies and academics, such as the Malaria Genomic Epidemiology Network (MalariaGEN) (www.malariagen.net/) and the Human Heredity and Health in Africa (H3Africa) program (h3africa.org/h3africa_whitepaper.pdf). These multi-national consortia have started to build the capacity in research skills and overcome the barriers for the use of genomics to address the disease burden of Africans. The time has come for African governments to heed the call of the World Health Organization to embrace genomics for the benefit of their populations.

The problems

There is a danger that new developments in genomics will increase the disparity in healthcare within countries, and between developed and developing countries, due to the high cost of implementing research findings in the clinic [1]. Inequalities in healthcare may be further exacerbated by the current intellectual property regime unless care is taken to ensure that knowledge produced through genomics research becomes a public good [3]. Furthermore, the limited biotechnology and information technology infrastructure in many African countries hinders the participation of researchers and clinicians in genomic medicine. There is a lack of market incentives for the private sector to invest in genomics research directed towards neglected diseases of the world’s poorest people. Therefore, the potential for genomics research to combat these diseases will only be realized through an enforceable global mechanism for public investment in health research on such diseases [3]. Indeed, more than one billion people in developing countries suffer from one or more neglected infectious diseases, most of which are found in Africa. Battling neglected infectious diseases is an important priority of the Bill & Melinda Gates Foundation (www.gatesfoundation.org), which has committed more than US$1.02 billion for research on methods of treatment and prevention, using genomic technologies in some cases.

Finally, the high cost of genomics and lack of technological capacity is compounded by a relatively low level of public and professional understanding of genomics in African countries [4]. There is, therefore, a need for African societies to prepare for the genomics era and its consequences through education at all levels, with particular emphasis on improving science education and the introduction of the principles of biomedical ethics to school children [1]. H3Africa draws increased attention to a number of longstanding and emerging issues in genetic and genomic research, such as informed consent, community engagement, privacy and confidentiality, use of genetic information, governance of biorepositories,
reciprocity/benefit sharing, and return of research results and incidental findings. These issues are highlighted by genome-wide association studies or whole exome/genome sequencing that will be carried out in many projects. Unfortunately, there is a dearth of published research on the views of Africans and African communities concerning participation in genomic research, and virtually no information on their perspectives regarding several of these issues [4].

The prospects
Simple clinical applications of DNA technology could provide immediate benefit for healthcare in many African countries. For example, sickle cell disease (SCD), an inherited condition, imposes a heavy economic burden on health systems in Africa, with nearly two-thirds of the 305,800 infants born annually with SCD worldwide living in Africa [5]. Prenatal genetic diagnosis of SCD would allow the option of termination of pregnancy. In Cameroon, despite the fact that the majority of doctors and parents of SCD children and adult SCD patients agreed with the principle of testing for prenatal genetic diagnosis of SCD (78.7%, 89.8% and 89.2%, respectively), parents (62.5% acceptance) were more in favor of termination of an SCD-affected pregnancy than doctors and adult patients (36.1% and 40.9% acceptance, respectively) [6]. These preliminary data signal potential value and adult patients (36.1% and 40.9% acceptance, respectively) 

The prospects for genomic medicine in Africa have been given a major boost by the US National Institutes of Health and the Wellcome Trust, which are investing approximately US$37 million in the H3Africa program over 5 years. The H3Africa consortium has three major goals: to increase the number of African scientists who are internationally competitive in genomics and population-based research; to establish collaborative networks of African investigators pursuing genomics-based disease-oriented projects on conditions that are prevalent in Africa; and to create and expand the infrastructure for genomics research, including bioinformatics and biorepositories on the African continent (http://h3africa.org/h3africa_whitepaper.pdf). This initiative, which was conceived by the African Society of Human Genetics, is set to expand the capacity of African institutions to undertake research in genomic medicine over the next decade.

A call to action
Improved capabilities of African institutions to undertake research will be realized if ministries of health implement at least three of the key recommendations of the World Health Organization Report on Genomics and World Health [1]. First, capacity building will be achieved most effectively through the development of academic public research and industrial partnerships between developed and developing countries and between developing countries themselves, encouraged by tax or other incentives, and through extensive networking within regions where there is evolving expertise in biotechnology. The H3Africa program has planted together researchers from 21 countries in Africa and elsewhere [10]. MalariaGEN has made major contributions to overcoming the scientific, ethical and practical challenges to carrying out such studies in Africa where the burden of malaria is greatest. The achievements of MalariaGEN include: analysis of 86,158 exonic single nucleotide polymorphisms in 227 samples, which has allowed the establishment of an open-access web application for the exploration of regional differences in allele frequency and of highly differentiated loci in the Plasmodium falciparum genome [10]; reviewing the methodological challenges of genome-wide association studies in Africa, and how these studies will be transformed by new approaches in statistical imputation and large-scale genome sequencing; the exploration of achieving valid consent both from the perspective of protecting individuals and building trust between communities and research groups, and recommendations of ethical data-release policies that provide adequate protection for the research aspirations of developing country scientists (http://www.sanger.ac.uk/research/initiatives/globalhealth/partnerships/malariagen.html).
the seed for capacity building, which needs to be nurtured by African governments to ensure the long-term sustainability of genomic medicine on the continent. Second, to be able to utilize the vast quantities of genomic data that are being generated, African countries must develop a critical mass of expertise in bioinformatics. There is already a shortage of trained scientists and technicians in this crucially important discipline, even in developed countries. As a first step, countries may decide to examine their current capacity in bioinformatics and identify existing shortfalls. Finally, all African countries need to evolve appropriate national frameworks to consider the ethical implications of genomics research and its applications in their own unique social, cultural, economic and religious context; empirical data from a broad spectrum of stakeholders and the public are needed for the development of effective genomics policies and programs of disease that affect the continent.

Abbreviations
H3Africa: Human Heredity and Health in Africa; HbF: fetal hemoglobin; MalariaGEN: Malaria Genomic Epidemiology Network; SCD: sickle cell disease.

Competing interests
The authors declare that they have no competing interests.

Published: 24 February 2014

References
1. World Health Organization: Genomics and World Health: Report of the Advisory Committee on Health Research. Geneva: World Health Organization; 2002.
2. Feero WG, Guttmacher AE, Collins FS. Genomic medicine - an updated primer. N Engl J Med 2001–2011, 2010:362.
3. Rottingen JA, Chamas C, Goyal LC, Harb H, Lagrada L, Mayosi BM. Securing the public good of health research and development for developing countries. Bull World Health Organ 2012, 90:398–400.
4. Wonkam A, Njamnshi AK, Angwafo FF 3rd. Knowledge and attitudes concerning medical genetics amongst physicians and medical students in Cameroon (sub-Saharan Africa). Genet Med 2006, 8:331–338.
5. Piel FB, Patil AP, Howes RE, Nyangiri OA, Gething PW, Dewi M, Temperley WH, Williams TN, Weatherall DJ, Hay SI. Global epidemiology of sickle haemoglobin in neonates: A contemporary geostatistical model-based map and population estimates. Lancet 2013, 381:142–151.
6. Wonkam A, de Vries J, Royal CD, Ramesar R, Angwafo FF. 3rd. Would you terminate a pregnancy affected by sickle cell disease? Analysis of views of patients in Cameroon. J Med Ethics 2013. doi:10.1136/medethics-2013-101392.
7. Makani J, Menzel S, Neya S, Cox SE, Drasar E, Soka D, Komba AN, Mgaya J, Rooks H, Vasavda N, Fegan G, Newton CR, Farall M, Thein SL. Genetics of fetal hemoglobin in Tanzanian and British patients with sickle cell anemia. Blood 2011, 117:1390–1392.
8. Mercier S, Küry S, Shaboodien G, Houniet DT, Khumalo NP, Bou-Hanna C, Bodak N, Cormier-Daire V, David A, Favre L, Figarella-Branger D, Gherardi RK, Glen E, Hamel A, Laboisse C, Le Caignec C, Lindenbaum P, Magot A, Munnich A, Mussini JM, Pillay K, Rahman T, Redon R, Salort-Campana E, Sambhanzoo-Koree M, Thavvin C, Barbarot S, Keavney B, Bézieau S, Mayosi BM. Mutations in FAM111B cause hereditary fibrosing poikiloderma with tendon contracture, myopathy, and pulmonary fibrosis. Am J Hum Genet 2013, 93:1100–1107.
9. Tekola Ayele F, Adeyemo A, Finan C, Hallu E, Sinnott P, Burlinson ND, Aseffa A, Rotimi CN, Newport MJ, Davey G. Hla class ii locus and susceptibility to podocytosis. N Engl J Med 2013, 366:1200–1208.
10. Manske M, Miotto O, Campino S, Aubeem S, Almagro-Garcia J, Maslen G, O’Brien J, Djimde A, Doumbo O, Zongo I, Ouedraogo JB, Michon P, Mueller I, Siba P, Nzila A, Bormann S, Kiara SM, Marsh K, Jiang H, Su XZ, Amaratunga C, Fairhurst R, Socheat D, Nosten F, Imwong M, White NJ, Sanders M, Anastasi A, Alcock D, Drury E, et al. Analysis of Plasmodium falciparum diversity in natural infections by deep sequencing. Nature 2012, 487:375–379.

doi:10.1186/gm528
Cite this article as: Wonkam and Mayosi: Genomic medicine in Africa: promise, problems and prospects. Genome Medicine 2014 6:11.