CURRICULUM VITA of Dr. Ruiqiang Li

Education

2010.11   Ph.D. in Biology, University of Copenhagen, Denmark

2002.06   B.S. in Physics, Southeast University, Nanjing, China.

Positions:

2011.03 –   Founder & CEO, Novogene, China

2009.07 – 2011.03   Vice President, BGI, China

2009.07 – 2011.03   Head of Research & Cooperation Division, BGI, China

2007.07 – 2009.07   Head of Bioinformatics Center, BGI, China

2004.09 – 2004.11   Visiting Scholar, Wellcome Trust Sanger Institute, Cambridge, UK

2002.07 – 2005.07   Group Leader, Bioinformatics Department, Beijing Genomics Institute (BGI)

Honorary Positions:

2009.12 -   Honorary Assistant Professor, Department of Computer Science, University of Hongkong, HK

Publications (Peer reviewed) (* Equal first author, # corresponding)

1. The Brassica rapa Genome Sequencing Project Consortium. (2011) "The genome of the mesoployploid crop species Brassica rapa." **Nature Genetics.** 43:1035-1039. doi:10.1038/ng.919.

2. Stukenbrock, E., Bataillon, T., Dutheil, J. Y., Hansen, T.T., **Li, R.**, Zala, M., et. al. (2011) "The making of a new pathogen: Insights from comparative population genomics of the domesticated wheat pathogen Mycosphaerella graminicola and its wild sister species." **Genome Res.** doi: 10.1101/gr.118851.110

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4. The Potato Genome Sequencing Consortium. (2011) "Genome sequence and analysis of the tuber crop potato." **Nature.** 475(7355):189-95. doi: 10.1038/nature10158. [cover story]

5. Li, Y., Zheng, H., Luo, R., Wu, H., Zhu, H., **Li, R.**, Cao, H. et. al. (2011) "Single-nucleotide resolution structural variations of two human genomes mapped by whole genome de novo assemblies." **Nature Biotech**. doi:10.1038/nbt.1904

6. Liu, Y., Gao, M., Lv, Y., …, **Li, R.** [#], Zhang, X. (2011) "Confirmation by exome sequencing of the pathogenic role of NCSTN mutations in Acne Inversa (Hidradenitis Suppurativa)." **Journal of Investigative Dermatology**. doi: 10.1038/jid.2011.62

7. Liaset, B., Hao, Q., …, **Li, R.**, et. al. (2011) "Nutritional Regulation of Bile Acid Metabolism Improves Pathological Characteristics of the Metabolic Syndrome." **The Journal of Biological Chemistry.** doi: 10.1074/jbc.M111.234732

8. Huang, S., Zhang, J., **Li, R.**, Zhang, W., He, Z., Lam, T., Peng, Z. and Yiu, S. (2011). "SOAPsplice: genome-wide ab initio detection of splice junctions from RNA-Seq data." **Front. Gene.** 2:46. doi: 10.3389/fgene.2011.00046

9. Shi, Y., Li, Y., …, **Li, R.**, et. al., (2011) "Exome sequencing identifies ZNF644 mutations in high myopia." **PLoS Genetics. 7:** e1002084.

10. Leung, C., Yiu, S., Yang, B., Peng, Y., …, **Li, R.**, Chin, F. (2011) "A robust and accurate binning algorithm for metagenomic sequences with arbitrary species abundance ratio." **Bioinformatics**. doi: 10.1093/bioinformatics/btr186

11. Korbel, J., Mills, R., … **Li, R.**, et. al. (2011) "Mapping copy number variation by population-scale genome sequencing." **Nature**. **470:** 59-65.

12. Wang, J., Yang, X., Xia, K., … Wang, J., **Li, R.**[#], Tang, B. (2010) "*TGM6* indentified as a novel causative gene of spinocerebellar ataxias using exome sequencing." **Brain**. doi:10.1093/brain/awq323

13. Wang, J., Zhang, J., **Li, R. ***, Zheng, H., et. al. (2010) "Evolutionary transients in the rice transcriptome." **Genomics, Proteomics & Bioinformatics.** doi:10.1016/S1672-0229(10)60023-X

14. Yang, B., Peng, Y., Leung, H., Yiu, S., Qin, J., **Li, R.**, Chin, F. (2010)

"MetaCluster: unsupervised binning of environmental genomic fragments and taxonomic annotation." **ACM-BCB.** 2010

15. Sudmant, P., Kitzman, J., Antonacci, F., Alkan, C., Malig, M., Tsalenko, A., Sampas, N., Bruhn, L., Shendure, J.; 1000 Genomes Project, Eichler, EE. (2010) "Diversity of human copy number variation and multicopy genes." **Science. 330(6004)**:641-6.

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21. Yi, X., Liang, Y., … **Li, R.**, Li, S., Yang, H., Nielsen, R., Wang, J., Wang, J. (2010) "Sequencing of 50 human exomes reveals adaptation to high altitude." **Science. 329**: 75-78.

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23. Zhang, G., Guo, G., Hu, X., Zhang, Y., Li, Q., **Li, R.**, Zhuang, R., et al. (2010) "Deep RNA sequencing at single base-pair resolution reveals high complexity of the rice transcriptome." **Genome Research**. **20**: 646-654.

24. **Li, R.***, Li, Y., Zheng, H., et al. (2010) "Building the sequence map of the human pan-genome." **Nature Biotech**. **28**: 57-63.

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27. Duan, J., Xia, Q., **Li, R.***, et al. (2010) "SilkDB v2.0: a platform for silkworm (*Bombyx mori*) genome biology." **Nucleic Acids Res. 38**: D453-456. (Database issue).

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30. Xia, Q., Guo, Y., … **Li, R.**, et al. (2009) "Complete resequencing of 40 genomes reveals domestication events and genes in silkworm (Bombyx)." **Science**. **326**: 433-436.

31. Huang S., **Li, R.***, et al. (2009) "The genome of the cucumber *Cucumis sativus sativus* L.." **Nature genetics**. **41**: 1275-1281. [cover story]

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