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Selected Scholarly Contributions [Data Provided by **SCOPUS**]

[Genome-wide association study provides evidence for a breast cancer risk locus at 6q22.33](#)

*Proceedings of the National Academy of Sciences of the United States of America*, 105 (11), pp. 4340-4345, 2008.

[Analysis of genetic variation in Ashkenazi Jews by high density SNP genotyping](#)

*BMC Genetics*, 9, art. no. 14, 2008.

[Power analysis for genome-wide association studies](#)

*BMC Genetics*, 8, art. no. 58, 2007.

[Expansion of symmetric exon-bordering domains does not explain evolution of lineage specific genes in mammals](#)

*Genetica*, 131 (1), pp. 59-68, 2007.

[Strong association of the Y402H variant in complement factor H at 1q32 with susceptibility to age-related macular degeneration](#)

*American Journal of Human Genetics*, 77 (1), pp. 149-153, 2005.

[Complement factor H polymorphism in age-related macular degeneration](#)

*Science*, 308 (5720), pp. 385-389, 2005.

[Estimating rates of alternative splicing in mammals and invertebrates \(multiple letters\)](#)

*Nature Genetics*, 36 (9), pp. 915-917, 2004.

[RSEARCH: Finding homologs of single structured RNA sequences](#)

*BMC Bioinformatics*, 4, art. no. 44, 2003.

[Noncoding RNA genes identified in AT-rich hyperthermophiles](#)

*Proceedings of the National Academy of Sciences of the United States of America*, 99 (11), pp. 7542-7547, 2002.

[Computational identification of noncoding RNAs in E. coli by comparative genomics](#)

*Current Biology*, 11 (17), pp. 1369-1373, 2001.