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Non-coding RNAs

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Deadline for manuscript
submissions:

closed (31 October 2017)

Message from the Guest Editors

Dear Colleagues,

Non-coding RNA has gained enormous interest in biomedical research during recent years. More and more researchers are starting to realize that the answers to some diseases may lie outside the coding regions. The National Human Genome Research Institute (NHGRI) launched the ENCODE consortium to study the non-coding regulatory genome, and the consortium claims that 80% of the human genome is associated with some kind of biochemical function through the regulation of the expression of coding genes. Nonetheless, current high throughput technology such as RNA-seq provides us with the opportunity to study non-coding RNA at an unprecedented level. We invite investigators to contribute original research articles, as well as review articles, that will stimulate the continuing efforts in the research of non-coding RNAs. Manuscripts submitted to this special issue are guaranteed to have a quick and fair review process.

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Guest Editors



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Special Issue



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Message from the Editor-in-Chief

Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised.

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