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EDITORIAL NOTE

The role of RNA in genetic programming, the interaction between humming bird and angiosperms in two areas of the Serra da Bodoquena (Mato Grosso do Sul, Bra and new extinct flying reptiles from China

ALEXANDER W.A. KELLNER Editor-in-chief

This issue of the *Anais da Academia Brasileira de Ciências* (AABC) is very special since, for the first tir journal has published over 1000 pages in a year. There are 29 articles that cover different scientific discipline mathematical sciences to vertebrate paleontology, providing important advances in these fields.

Among the highlights of the present issue is the contribution of John S. Mattick (University of Queensland bane, Australia) regarding genetic programming. For the last 50 years or so, there has been a general belief most of the genetic information for multicellular organisms is transacted by proteins. The starting point assumption is the general 'DNA makes RNA makes protein' dogma (e.g., Mattick 2010). Although this might for simple unicellular prokaryotes, there is growing evidence that RNA plays a more central role in the gene gramming system for eukaryotes. The present approach is that the genomes of complex organisms, here in humans, can be regarded as islands of protein-coding sequences (e.g., Ovcharenko et al. 2005), with RNA cout the major role regarding regulation. As observed previously, this molecule can form rather sophisticated tures among others) whose study – as Mattick (2010) points out – might hold the key to understand human even with far more outreaching possibilities than previously thought, including the comprehension of individual subilities to complex diseases.

As has been pointed out several times in the literature, phenology and pollination are paramount in the reprod of plants. While phenology is concerned with a variety of factors such as interactions between plants and a including annual cycles and seasonal rhythms of individual plant species (e.g., Talora and Morellato 2000), ro on pollination focus on factors such as the amount of mutual dependence among species that encompasses source and supply for pollen vectors (e.g., Smith-Ramírez et al. 2005). In an interesting study published issue of the AABC, Rogério R. Faria and Andréa C. Araújo from the *Universidade Federal do Mato Gros Sul* (Campo Grande) discuss the interaction between hummingbirds (which are considered among the most et pollinators, particularly in the neotropics) and bird-pollinated plants in two areas of the Serra da Bodoquena sin the mid-south region of Mato Grosso do Sul. This type of study might help us to better understand the of deforestation in the local biota, particularly in Mato Grosso do Sul State, where agriculture is expanding

in the last decades at the expense of the indigenous forests.



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in the Yixian and Jiufotang Formations (e.g., Zhou et al. 2003). Most recently, however, several new specimens have also been unearthed in stratigraphic units that are older and that are referred in the literature as the Daohugou or Tiaojishan Formations (Wang et al. 2010), particularly from a region called Linglongta (western Liaoning). Among these are remains of what is perhaps the most interesting clade of pterosaurs (an extinct group of flying reptiles) discovered in the last years: the Wukongopteridae. This group of pterosaurs, which was proposed in a paper published by the AABC (Wang et al. 2009) revealed a quite unusual combination of primitive and derived characters of a new species, *Wukongopterus lii*, which became even more apparent with the discovery of *Darwinopterus modularis* from the same deposits (Lü et al. 2010). Here, Xiaolin Wang (Institute of Vertebrate Paleontology and Paleoanthropology, Chinese Academy of Sciences, Beijing, China) and colleagues report two new species that can be attributed to the Wukongopteridae (Wang et al. 2010). In several color plates, these authors unravel the detailed anatomy of these flying reptiles that proved to be more diversified than previously known and surely will open a new chapter in pterosaur studies.

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