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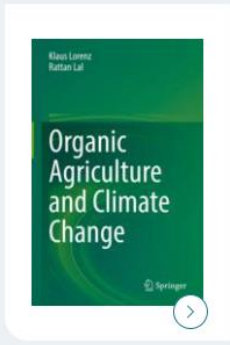
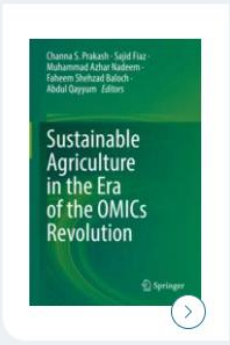
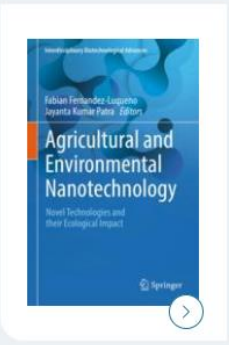
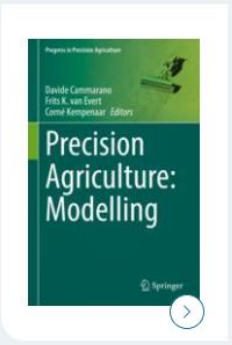
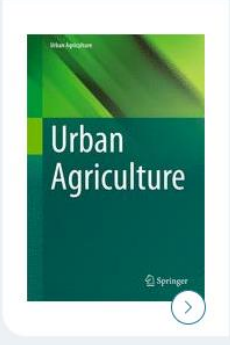
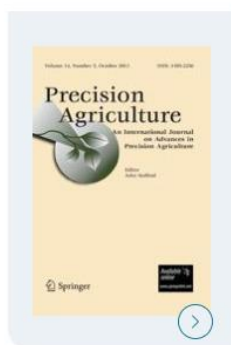
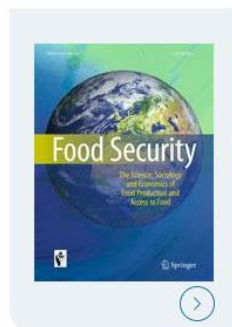
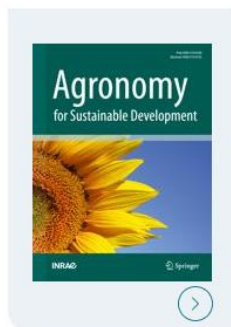
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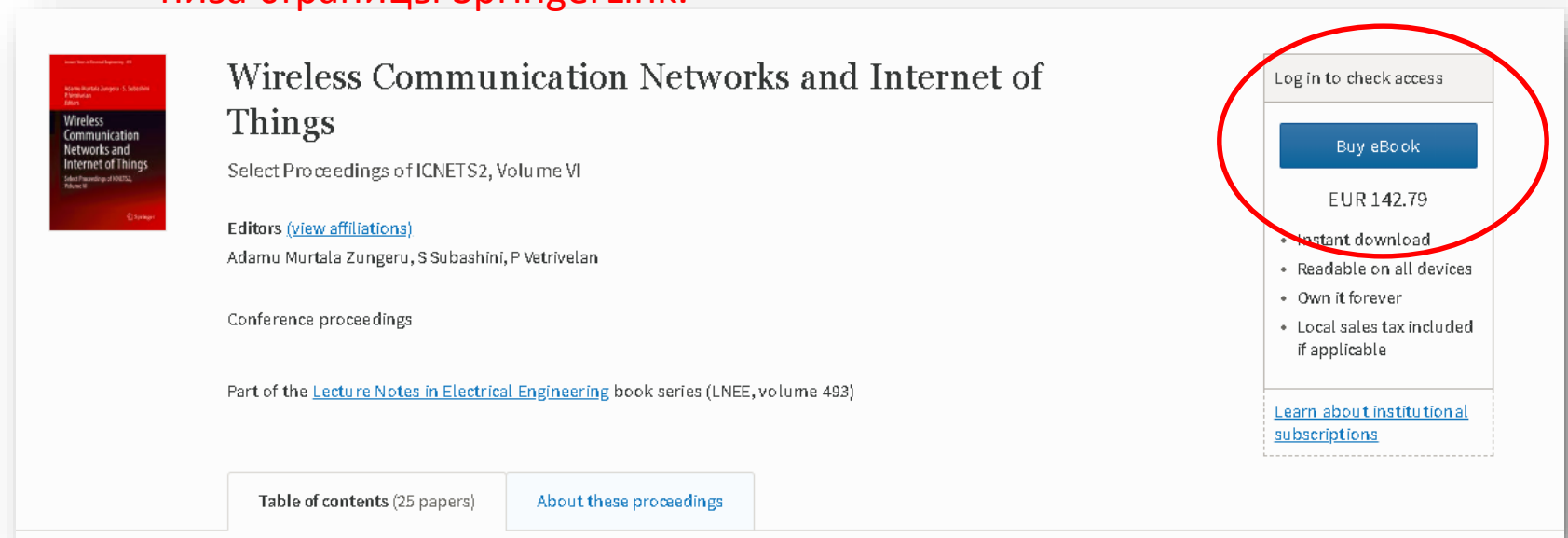
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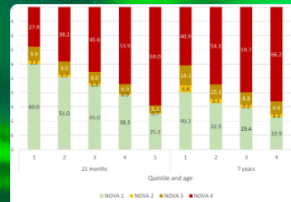
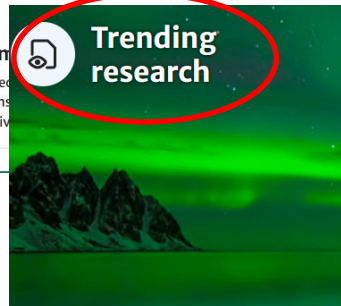
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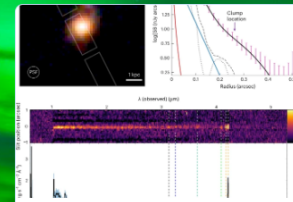
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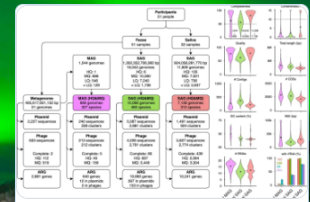
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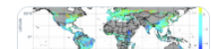
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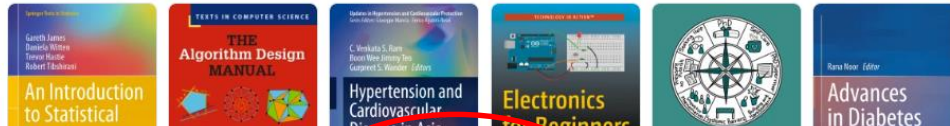
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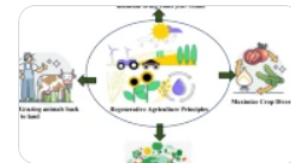
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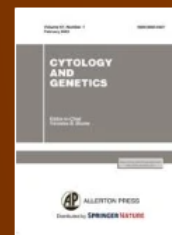
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Molecular Phylogeny and Genetic Diversity of Carpathian Members of the Genus *Muscari* Inferred from Plastid DNA Sequences (1)

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The genus *Muscari* Mill. (Asparagaceae Juss.) includes about 80 species distributed in Eurasia, mainly in the Mediterranean region. Recent molecular phylogenetic studies have shown that the taxa belonging to this group form a monophyletic clade and are closely related. However, the phylogeny and status of some taxa of *Muscari* sensu lato remain controversial. So far, most phylogenetic studies of the genus *Muscari* have used almost exclusively Mediterranean plant material, while representatives of the genus from the

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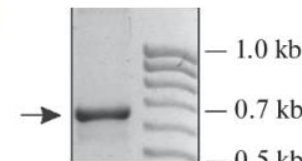


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This fully revised third edition includes up-to-date topics and developments in the field, which has made tremendous strides since the publication of the second edition in 2004. Many novel techniques based on Next Generation Sequencing have sped up the analysis of fungi and major advances have been made in genome editing, leading to a deeper understanding of the genetics underlying cellular processes as well as their applicability. At the same time, the relevance of fungi is unbroken, both due to the serious threats to human health and welfare posed by fungal pests and pathogens, and to the many benefits that fungal biotechnology can offer for diverse emerging markets and processes that form the basis of the modern bioeconomy.

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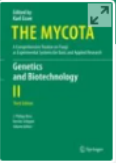
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