

Research

- 606 **In-depth Proteome of the Hypopharyngeal Glands of Honeybee Workers Reveals Highly Activated Protein and Energy Metabolism in Priming the Secretion of Royal Jelly**
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- 622 **Quantitative Mass Spectrometry to Interrogate Proteomic Heterogeneity in Metastatic Lung Adenocarcinoma and Validate a Novel Somatic Mutation CDK12-G879V**
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- 642 **High-density Peptide Arrays Help to Identify Linear Immunogenic B-cell Epitopes in Individuals Naturally Exposed to Malaria Infection**
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- 715 **Proteomics Reveals Multiple Phenotypes Associated with N-linked Glycosylation in *Campylobacter jejuni***
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On the cover: Antibodies are a fundamental part of the immunity to severe malaria, but they are not well characterized. Therefore, a high-throughput peptide array technology was used to analyze the antibodies of patients against epitopes of the *P. falciparum* proteome. Known and novel vaccine candidates can be quickly identified. This serves as a general approach to develop vaccines and biomarkers. For details, see the article by Loeffler *et al.*, pages 642–656.

- 735 **Identification of Candidate Plasma Protein Biomarkers for Cervical Cancer Using the Multiplex Proximity Extension Assay**
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