

Program

17:00 - 17:25 Katrin Kemmerer

AK Weigand, TU Darmstadt

Hypoxia reduces MAX expression in endothelial cells by unproductive splicing

17:25 - 17:50 Katja Sträßer

AK Sträßer, Uni Gießen

Genome-wide data reveals molecular functions of mRNA-binding proteins

17:50 - 18:00 Break

18:00 - 18:25 Jun Yang

AK Entian, Goethe-University

Ribonucleome analysis of yeast 18S rRNA

18:25 - 18:50 Kathi Zarnack

AK Zarnack, Goethe-University

Control of a neuronal morphology program by the RNA-binding zinc finger protein
Unkempt

Pizza & Drinks