

Program of the 20th TBI Winterseminar in Bled 2005

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|----|---|-----------------------|---|
| Mo | 1 | Michael Wolfinger | Pin(girl)fold |
| | 1 | Andreas Svrcek-Seiler | Fighting slow electrostatic Approximations |
| | 1 | Stephan Steigrale | Distinct Sets of Antisense Transcripts with Common Functions |
| | 2 | Peter Stadler | Molecular Phylogenies Without Aligned Sequences |
| | 2 | Matthias Kruspe | A Multiple Alignment Tool in 3D |
| Di | 1 | Claudia Fried | Single Nucleotide Polymorphisms in the genome |
| | 1 | Paul Perco | Genetic Algorithms for Genome Analysis |
| | 1 | Axel Mosig | Spicy Barbeque |
| | 1 | Sandro Wenzel | Multicanonical Monte Carlo Simulations |
| | 2 | Bärbel Stadler | Why relations are topologies |
| | 2 | Peter Stadler | Intro to Wilfried's Talk |
| | 2 | Wilfried Imrich | Prime Factor Theorem for a Generalized Strong Product |
| | 2 | Josef Leydold | A note on the Courant-Herrmann-Conjecture for manifolds |
| Mi | 1 | Tomo Pisanski | |
| | 1 | Andrea Tanzer | Micro RNAs: Regulators Regulating Regulators |
| | 1 | Roman Stocsits | Fantasy, Brain Damage, and My Dreams of Perfectly Simulating Genome Evolution |
| | 1 | Stefan Washietl | Tracking down functional RNAs in the human genome |
| | 2 | Kristin Missal | non-coding RNAs in Ciona sp. (part 2A) |
| | 2 | Dominic Rose | non-coding RNAs in Ciona sp. (part B1) |
| | 2 | Manja Lindemeyer | Evolution of snRNAs |
| Do | 1 | Ivo Hofacker | Folding Dynamics of RNA Secondary Structures |
| | 1 | Ulli Mückstein | <i>In silico</i> Study of miRNA binding-sites |
| | 1 | Stephan Bernhart | About RNAfold and it's variants |
| | 2 | Sonja Prohaska | Click to add title (by AutoContent Wizard) |
| | 2 | Lukas Endler | Mathematical Analysis of Cyclic Gene Regulatory Networks |
| Fr | 1 | Gil Benkő | Oranization Theory applied to Chemical Reaction Networks |
| | 1 | Christoph Flamm | Loos Ends, Remarks, Footnotes, ... |
| | 1 | Camille Stephan | The Amoeba-Flagellat Transformation |
| | 1 | Konstantin Klemm | From Las Vegas to Monte Carlo and back: Sampling cycles in graphs |
| | 1 | Dirk Drasto | Individual cell-based models of the spatio-temporal organisation of multicellular systems |

Session 1 from 17:00-19:00; Session 2 from 20:30-22:30