

Lucian Ilie



- Lucian Ilie, Professor
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Research – Bioinformatics

- Genomics

- Sequence Analysis

- Efficiently detecting similarities between DNA sequences (SpEED)
 - Whole genome alignment (E-MEM)
 - Read mapping (SHRiMP2)

- DNA Sequencing – Finding DNA sequences of real genomes

- Error correction (RACER)
 - De novo genome assembly (SAGE)
 - Assembly evaluation (LASER)
 - Illumina, Pacific Biosciences (HISEA)

- Probe design – Unique probes to identify, e.g., missing genes

- DNA oligonucleotide design (BOND)

- Proteomics

- Protein Protein Interactions (PPI)

- Predicting interactions between proteins (SPRINT)
 - Aligning networks of PPIs between different organisms

Lab

- Our own computer cluster
 - 1 node: 32 cores, 1TB RAM
 - 4 nodes: 12 cores, 256GB RAM each



Teaching

- CS9877 – Research Topics in Genomics and Proteomics
 - fall 2016
 - no background required
 - introduction to basic algorithms in sequence analysis
 - currently hot research topics
 - no assignments or exams (only presentations)
 - www.csd.uwo.ca/~ilie/courses/CS9877.html