



# Bayesian Network Tools in Java (BNJ) v2.0

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<http://bndev.sourceforge.net>



## What is BNJ?

- Software toolkit for research and development using graphical models
- Open source (GNU General Public License)
- 100% Java (J2EE v1.4)
- Developed at KDD Lab, Kansas State University
- <http://bndev.sourceforge.net>
- Version 2 currently in alpha stage



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# Intended Users

- Researchers / students
  - Experiment with algorithms for learning, inference
    - Standardized comparison
    - Synthesis
  - Create, edit, convert **networks, data sets**
- Developers
  - New algorithms for graphical models using BNJ API
  - Applications



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# BNJ History

- BNC: initiated 1997, U. Illinois
- BNJ 1: developed 1999-2002, KS State
  - Hard to maintain
  - Redesigned from scratch
- BNJ 2: development started Dec 2002
  - Surpasses BNJ v1 in features, flexibility, performance
  - More standardized API



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## BNJ Highlights [1]: Network Interchange

- 8 network formats supported
  - Hugin .net (both 5.7 and 6.0)
  - XML-Bif
  - Legacy BIF
  - Microsoft XBN
  - Legacy DSC
  - Genie DSL
  - Ergo ENT
  - LibB .net
- Opens, saves, converts



## BNJ Highlights [2]: Data Formats Supported

- Microsoft Excel (.xls)
- WEKA (.arff)
- LibB data
- XML-data
- Legacy .dat format
- Flat files
  - Space/tab delimited ASCII .txt
  - Comma-separated



## BNJ Highlights [3]: Exact Inference

- Junction Tree [Lauritzen & Spiegelhalter, 1988]
- Variable elimination [Shenoy; Dechter] with optimizations
  - *JavaBayes* [Cozman, 2001]
  - Kansas State KDD Lab [Joehanes & Hsu, 2003]
- Singly-connected network belief propagation [Pearl, 1983]
- Cutset Conditioning – under revision [Suermondt, Horvitz, & Cooper, 1990]



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## BNJ Highlights [4]: Approximate Inference

- Sampling based:
  - Logic Sampling
  - Forward Sampling
  - Likelihood Weighting
  - Self-Importance Sampling
  - Adaptive Importance Sampling (AIS)
- Bounded Cutset Conditioning (BCC) – under revision
- Hybrid: AIS-BCC bridge – under revision



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## BNJ Highlights [5]: Structure Learning

- Greedy (Bayesian Dirichlet) score-based: *K2* [Cooper & Herskovits, 1992]
- Genetic wrapper
  - cf. [Larranaga, 1998; Hsu, Guo, Perry, Stilson, 2002]
  - GAWK (for *K2*) [Joehanes, 2003]
  - Direct structure learning [Perry, 2003]
- Iterative Improvement
  - Straightforward hill-climbing
  - Simulated annealing (SA)
  - SA with adversarial reweighting
  - Other algorithms



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## BNJ Highlights [6]: Analysis and Experimentation

- Structure scoring during, after learning
  - Graph errors
  - RMSE
  - Log likelihood score
  - Dirichlet structure score
- Robustness analysis module
- Data generator: applies existing sampling-based inference algorithms



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## BNJ Highlights [7]: Probabilistic Relational Models

- Preliminary support for PRM structure learning
  - Accesses relational databases (*mySQL*, *PostgreSQL*, *ORACLE 9i*) via JDBC interface
  - Preliminary local database loading support (without any database engines)
  - Currently: adapt traditional learning algorithms such as *K2*, *Sparse Candidate*, etc. to relational models
- PRM inference: planned for full release of v2 (Spring, 2004)



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## BNJ Highlights [8]

- Converter Factory
  - Standalone application
  - GUI front-end
  - Converts among supported network, data formats
- Database GUI Tool
  - Transfer data files to and from server
  - Submit SQL commands through JDBC interface
  - Currently used for PRM learning



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## BNJ Highlights [9]

- Wizards for
  - Inference*
  - Learning*
  - Others planned
- GUI for Network Editing
  - Still in redevelopment
  - Currently display-mode only
- All tools available in command-line mode



## BNJ Performance

- Relatively fast inference for small to medium networks
- Tends to slow down when node arity high
- Optimization underway
- Very fast learning engine
  - 235 nodes, 76 data points (yeast cell-cycle expression data, Spellman-Gasch) with  $K2$ : 3 seconds on AMD Athlon XP 1.6GHz
  - Full alarm (37 nodes, 3000 data points) with  $K2$ : 13 seconds on AMD Athlon XP 1.6GHz



# Applications, New Research: What We Have Done with BNJ

- Computational genomics:  
learning gene expression pathways
  - *Saccharomyces cerevisiae* (yeast)  
[Johanes & Hsu, 2003]
  - *Oryza sativa* (rice) defense-response – in progress  
<http://www.kddresearch.org/REU/Summer-2003>
- PRM Learning Experiments: *EachMovie* data
- New Developments
  - Variable ordering wrappers [Hsu *et al.*, 2002]
  - Hybrid inference algorithms (AIS-BCC)



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# Software Demo

- Development using Eclipse platform
  - Open-source IDE
  - From IBM ([www.eclipse.org](http://www.eclipse.org))
- Standalone applications: *coming soon*
- Sources, documentation on *SourceForge*  
<http://bndev.sourceforge.net>



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