



Search methods exploiting the problem structure

Tutorial – Kiev 09/2010

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R. Marinescu (4C, Ireland), P. Jeavons (Univ. Oxford, UK), and M. Cooper (IRIT, France)

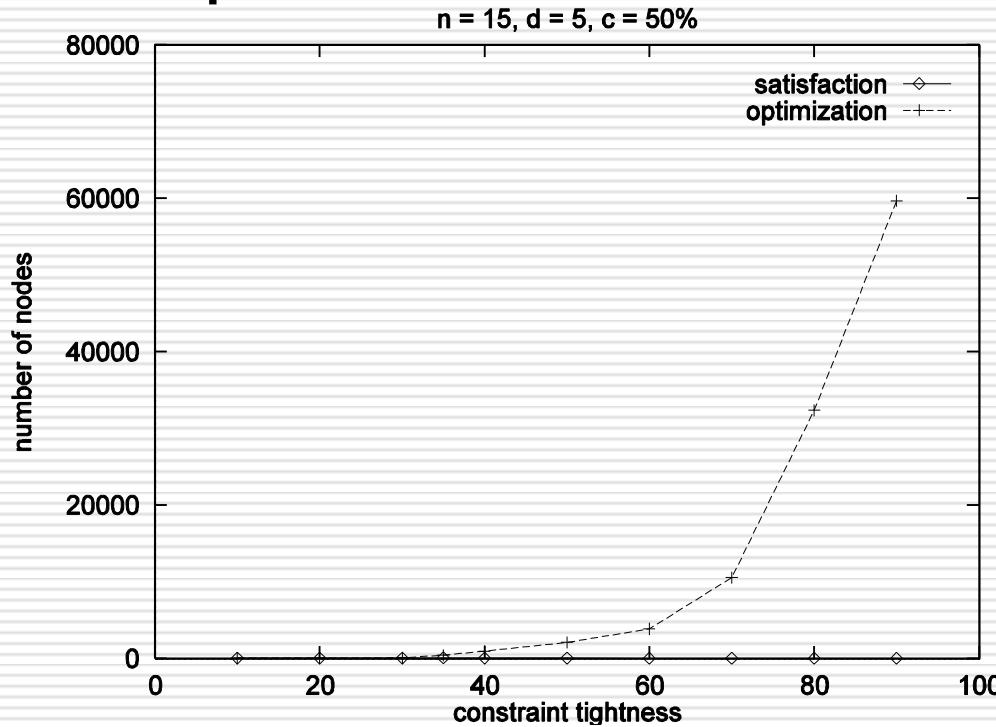
- (X, D, F, k)
 - $X = \{X_1, \dots, X_n\}$ **n** variables
 - $D = \{D_1, \dots, D_n\}$ **n** finite domains of maximum size **d**
 - $F = \{f_1, \dots, f_m\}$, a set of **m** cost functions
 - ★ f_S with scope S , assigns a positive cost to any assignment of S
 - ★ f_\emptyset problem lower bound (obtained by enforcing local consistency by *Equivalence Preserving Transformations*)
 - **k**, maximum cost and global upper bound
- Find an assignment A of all the variables which **minimizes**

$$\sum_{f_S \in F} f_S (A[S])$$

NP-hard problem

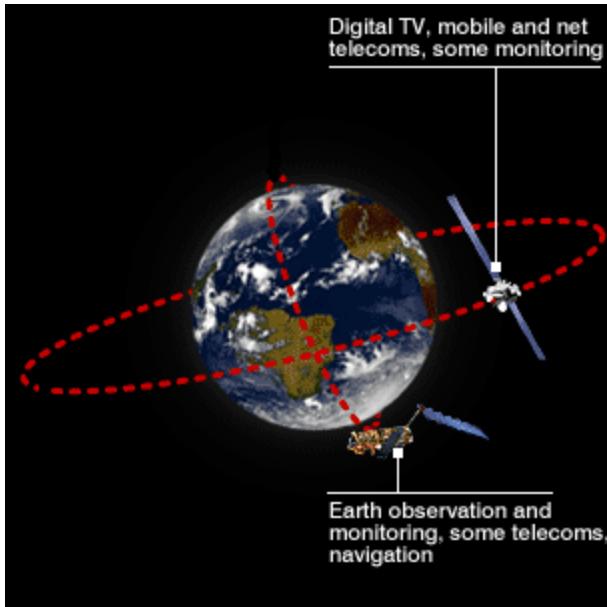
Observation

- Optimization problems are harder than satisfaction problems



CSP vs. Max-CSP

Combinatorial Optimization



Find a schedule for the satellite that **maximizes** the number of photographs taken, subject to the on-board recording capacity

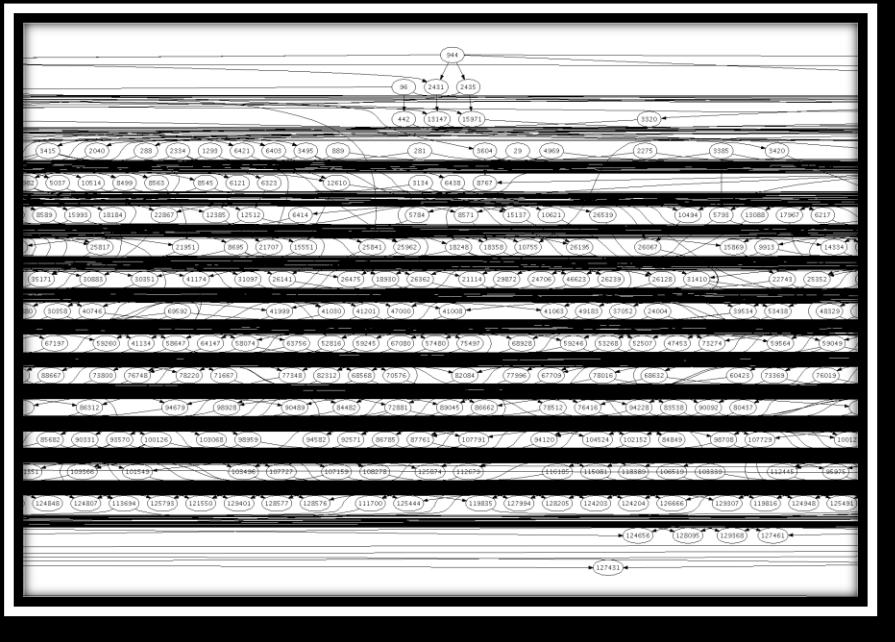
$$n \leq 364, e \leq 10,108$$



Assign frequencies to a set of radio links such that interferences are **minimized**

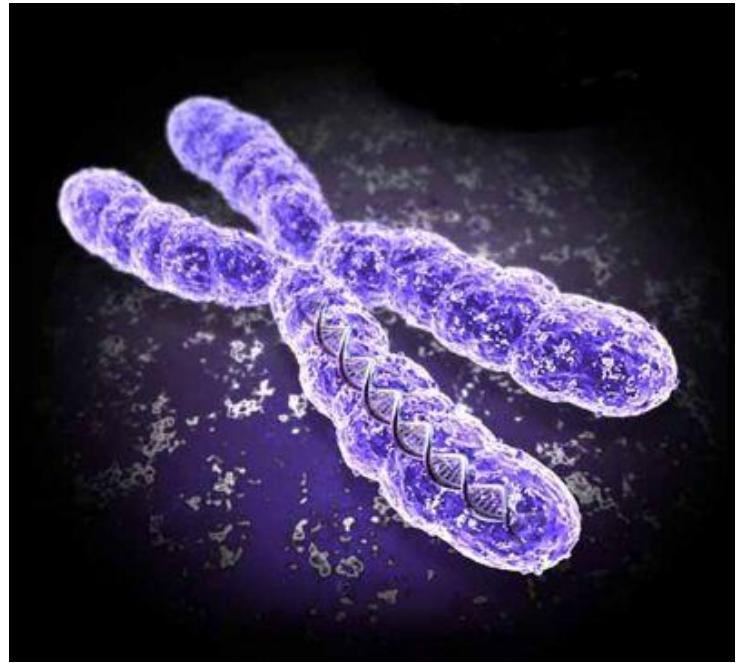
$$n \leq 458, e \leq 5,000$$

Combinatorial Optimization



Find a **minimum** number of erroneous genotypes such that their removal restores consistency in a complex pedigree

$$n \leq 20,000, e \leq 30,000$$

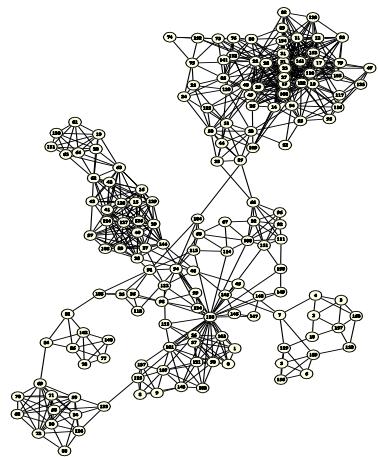


Select a **minimum** number of markers such that they cover all the markers and secondly, **maximize** weighted coverage sum of unselected markers and **maximize** dispersion between selected markers

$$n \leq 1,500, e \leq 150,000$$

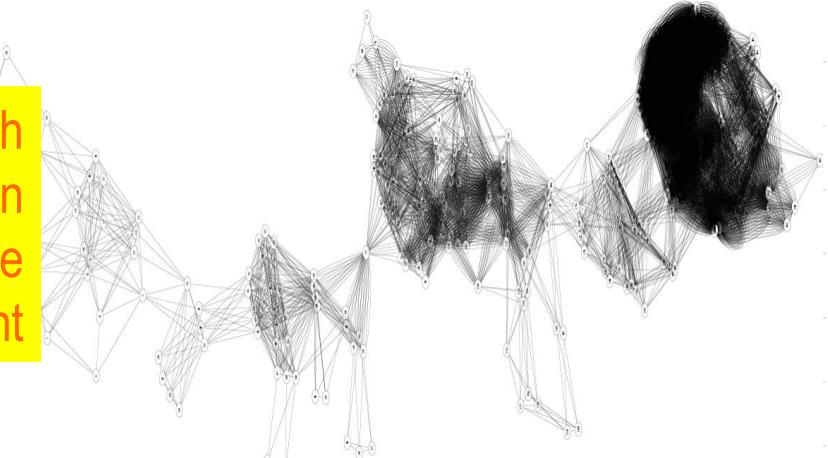
Many real applications have a structured network

Radio
Link
Frequency
Assignment



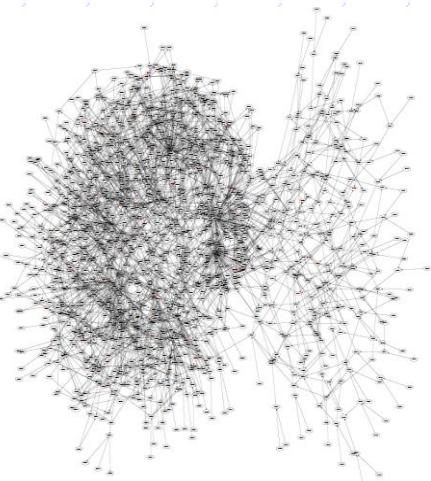
CELAR SCEN-07r
(*Constraints* 4(1), 1999)

Earth
Observation
Satellite
Management



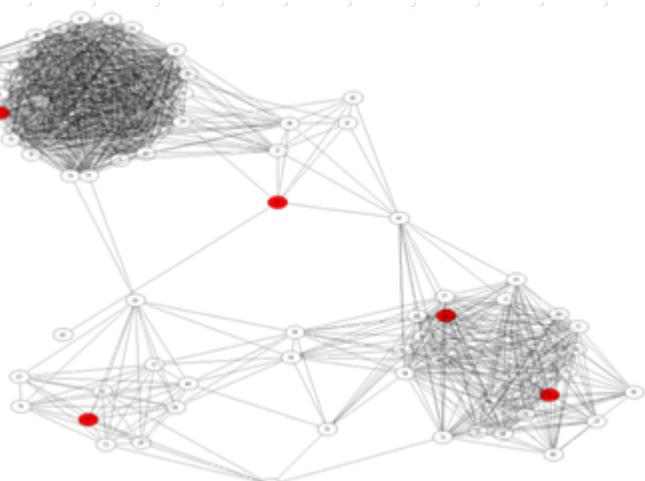
SPOT5 #509 (*Constraints* 4(3), 1999)

Mendelian
Error
Detection



langladeM7 sheep pedigree
(*Constraints* 13(1), 2008)

Tag
SNP
Selection



HapMap chr01 $r^2 \geq 0.8$ #14481
(*Bioinformatics* 22(2), 2006)

CELAR 6 results since 1993

n. of vars: n=100, domain size: d=44, n. of cost functions: e=1222

Time of optimality proof	Method(s) used	Publication	
26 days (SUN UltraSparc 167 MHz)	Ad-hoc problem decomposition & Russian Doll Search (<i>22 vars only</i>)	(de Givry, Verfaillie, Schiex, CP 1997)	
3 days (SUN Sparc 2)	Ad-hoc problem decomposition & PFC-MRDAC (<i>22 vars only</i>)	(Larrosa, Meseguer, Schiex, AIJ 1998)	
8 hours (DEC Alpha 500MP)	Preprocessing rules & Cluster Tree Elimination	(Koster PhD thesis, 1999)	
3 hours (PC 2.4 GHz)	B&B with EDAC & tree decomposition (BTD)	(de Givry, Schiex, Verfaillie, AAAI 2006)	
1' 26" (PC 2.5 GHz)	25000x 16 x	BTD & variable ordering heuristic & dichotomous branching	(Sanchez, Allouche, de Givry, Schiex, IJCAI 2009)

CELAR 7 (n=200) solved in 4.5 days (Sanchez et al, IJCAI 2009)

CELAR 8 (n=458) solved in 127 days (Allouche et al, CP 2010)

All CELAR and GRAPH instances are closed!

Solving methods

BTD, AND/OR graph search

Time: $\exp(\text{treewidth})$

Space: $\exp(\text{treewidth})$

Search: Conditioning

Time: $\exp(n)$

Space: linear

BB-VE

Time: $\exp(n)$

Space: linear

Inference: Elimination

Time: $\exp(\text{treewidth})$

Space: $\exp(\text{treewidth})$

Complete

Depth-First
Branch & Bound

A*

Incomplete

Simulated Annealing
Tabu Search, VNS,
Stochastic Local Search

Incomplete

Local Consistency

Mini-bucket(i)

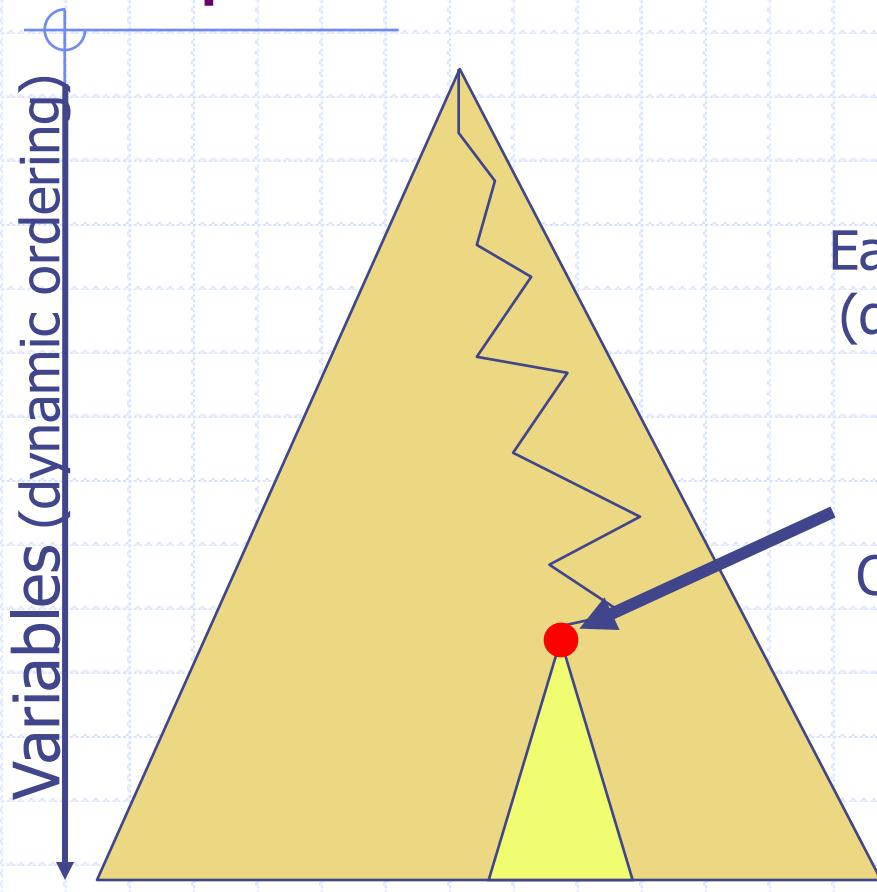
Hybrids

Complete

Variable Elimination

Cluster Tree Elimination

Depth-First Branch and Bound (DFBB)



Each node is a WCSP subproblem
(defined by current conditioning)

(LB) Lower Bound = f_{\emptyset}
under estimation of the
best solution
in the sub-tree

If $LB \geq k$ then prune

(UB) Upper Bound
= best solution found so far = k

Combination of cost functions

A	B	f(A,B)
b	b	6
b	g	0
g	b	0
g	g	6

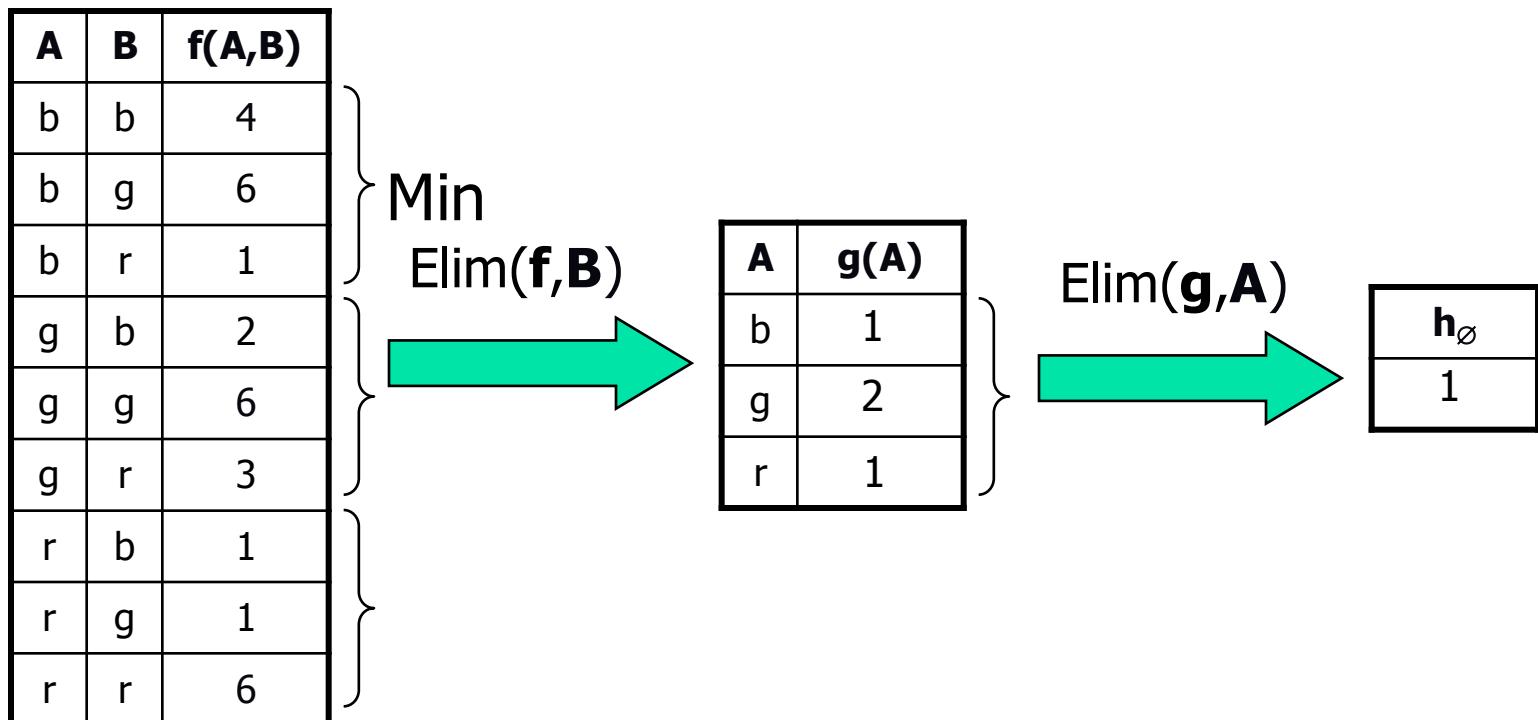


A	B	C	f(A,B,C)
b	b	b	12
b	b	g	6
b	g	b	0
b	g	g	6
g	b	b	6
g	b	g	0
g	g	b	6
g	g	g	12

B	C	f(B,C)
b	b	6
b	g	0
g	b	0
g	g	6

$$= 0 + 6$$

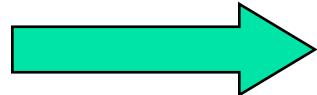
Elimination in a cost function



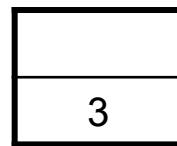
Conditioning a cost function

A	B	$f(A,B)$
b	b	6
b	g	0
b	r	3
g	b	0
g	g	6
g	r	0
r	b	0
r	g	0
r	r	6

Assign(\mathbf{f}_{AB}, A, b)



$g(B)$



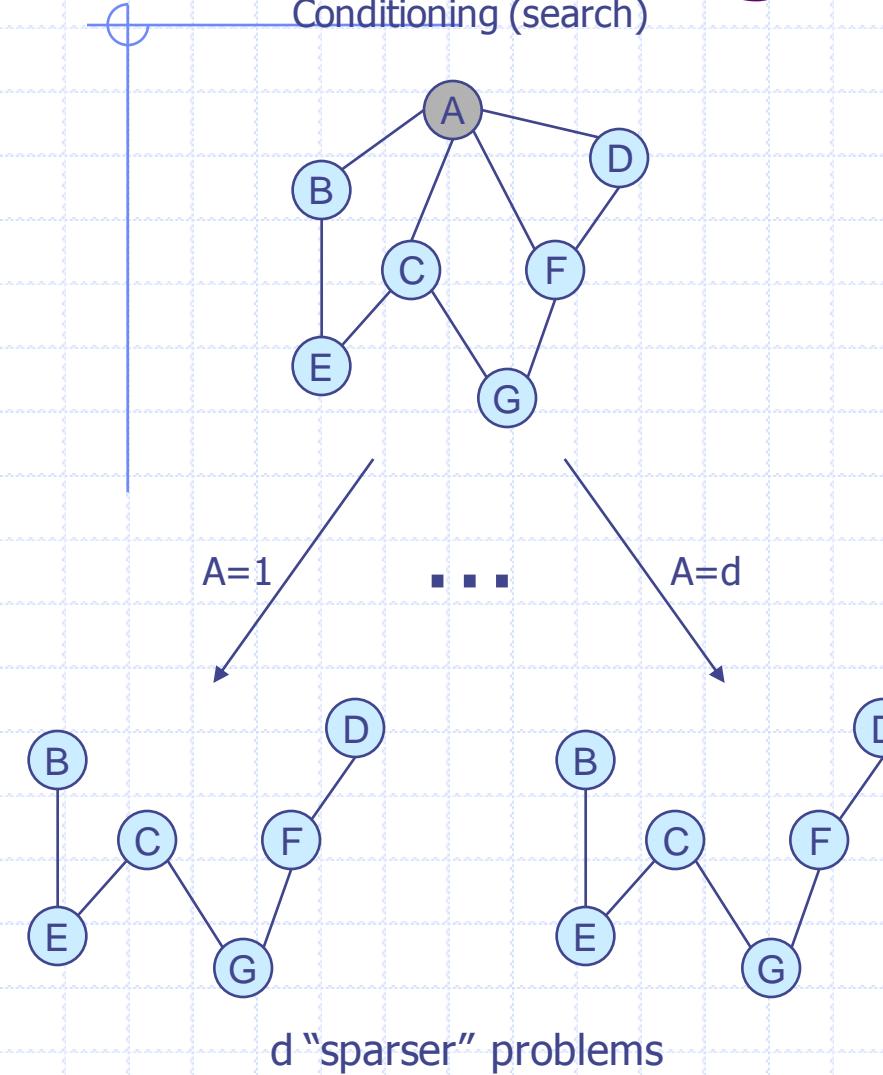
Assign(\mathbf{g}, B, r)



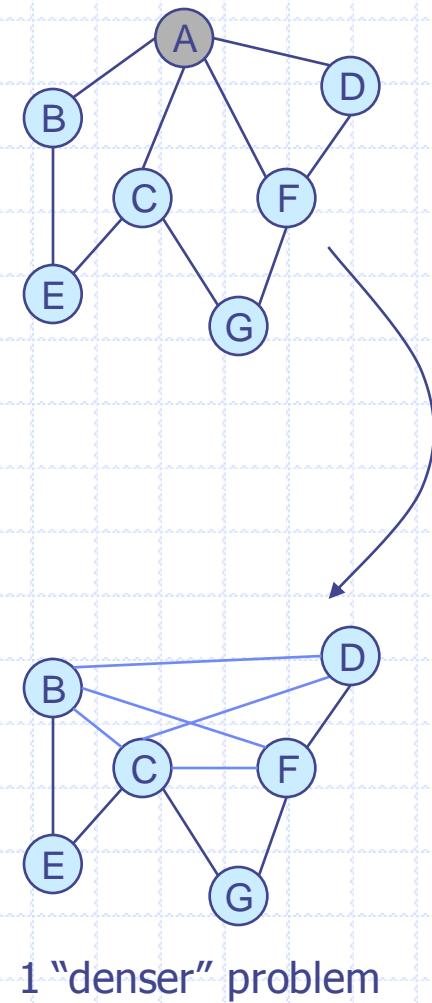
h_\emptyset

Conditioning vs. Elimination

Conditioning (search)



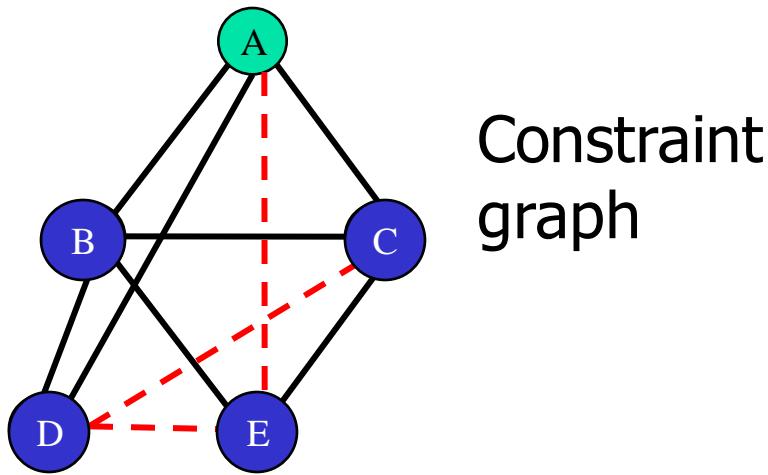
Elimination (inference)



Variable elimination (aka bucket elimination)

- Eliminate Variables one by one.
 - When all variables have been eliminated,
the **problem is solved**
 - Optimal solutions of the original problem
can be recomputed
-
- Complexity: exponential in the *induced width*
(maximum degree found during the elimination process)
 - Minimizing induced width is NP-hard.

Computing the Optimal Cost Solution



$$\text{OPT} = \min_{e=0,d,c,b} f(a,b) + f(a,c) + f(a,d) + f(b,c) + f(b,d) + f(b,e) + f(c,e)$$

Combination

$$\min_{e=0} \min_d f(a,d) + \min_c f(a,c) + f(c,e) + \min_b f(a,b) + f(b,c) + f(b,d) + f(b,e)$$

Variable Elimination

$$h^B(a, d, c, e)$$

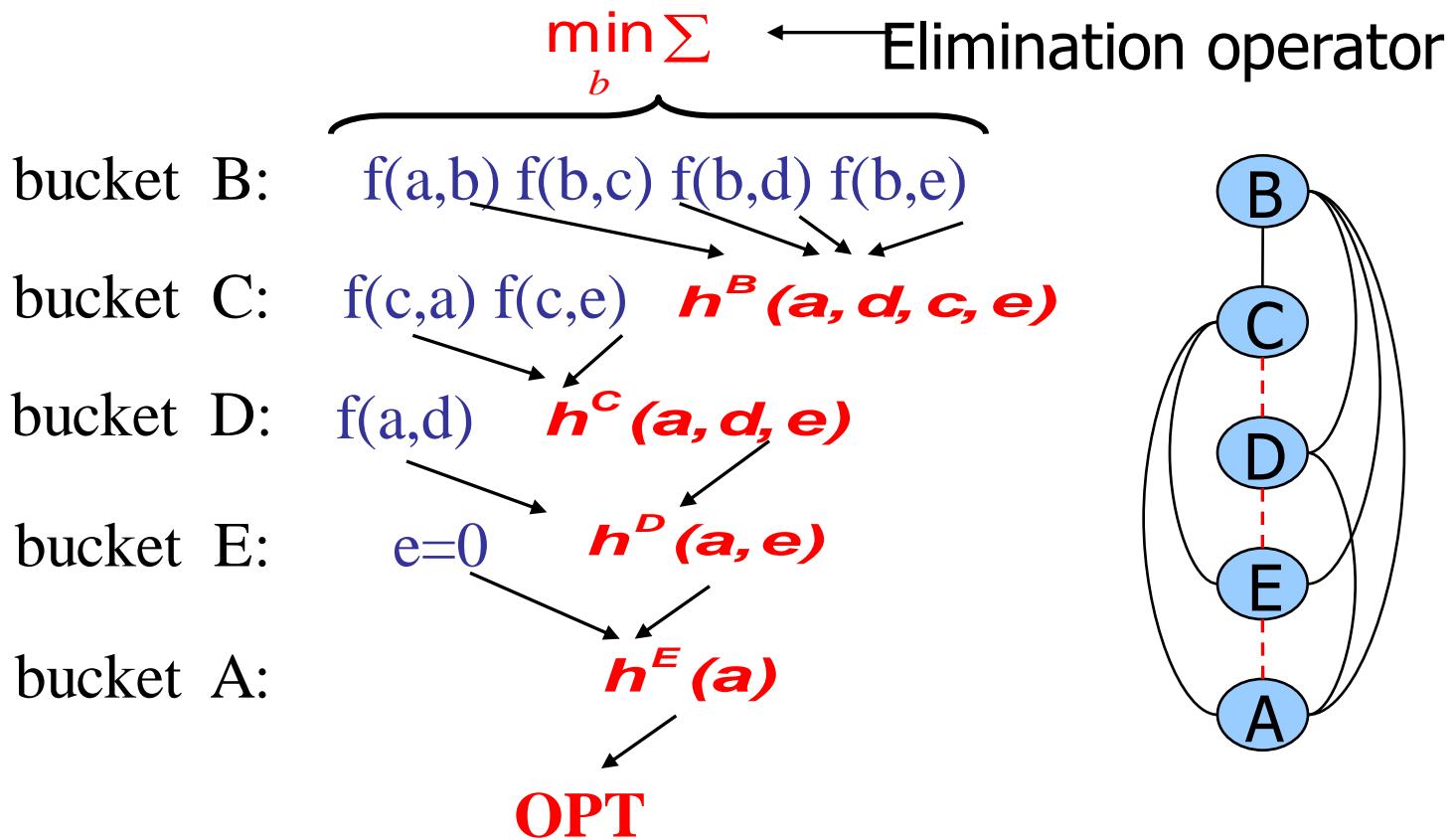
Finding

$$OPT = \min_{X_1, \dots, X_n} \sum_{j=1}^r f_j(X)$$

Algorithm **elim-opt** (Dechter, 1996)

Non-serial Dynamic Programming (Bertele & Briochi, 1973)

$$OPT = \min_{a,e,d,c,b} F(a,b) + F(a,c) + F(a,d) + F(b,c) + F(b,d) + F(b,e) + F(c,e)$$



Generating the Optimal Assignment

$$5. \ b' = \arg \min_b f(a', b) + f(b, c') +$$

$$+ f(b, d') + f(b, e')$$

$$4. \ c' = \arg \min_c f(c, a') + f(c, e') +$$

$$+ h^B(a', d', c, e')$$

$$3. \ d' = \arg \min_d f(a', d) + h^C(a', d, e')$$

$$2. \ e' = 0$$

$$1. \ a' = \arg \min_a h^E(a)$$



$$B: \quad f(a, b) \ f(b, c) \ f(b, d) \ f(b, e)$$

$$C: \quad f(c, a) \ f(c, e) \quad \quad h^B(a, d, c, e)$$

$$D: \quad f(a, d) \quad \quad h^C(a, d, e)$$

$$E: \quad e=0 \quad \quad h^D(a, e)$$

$$A: \quad \quad \quad h^E(a)$$

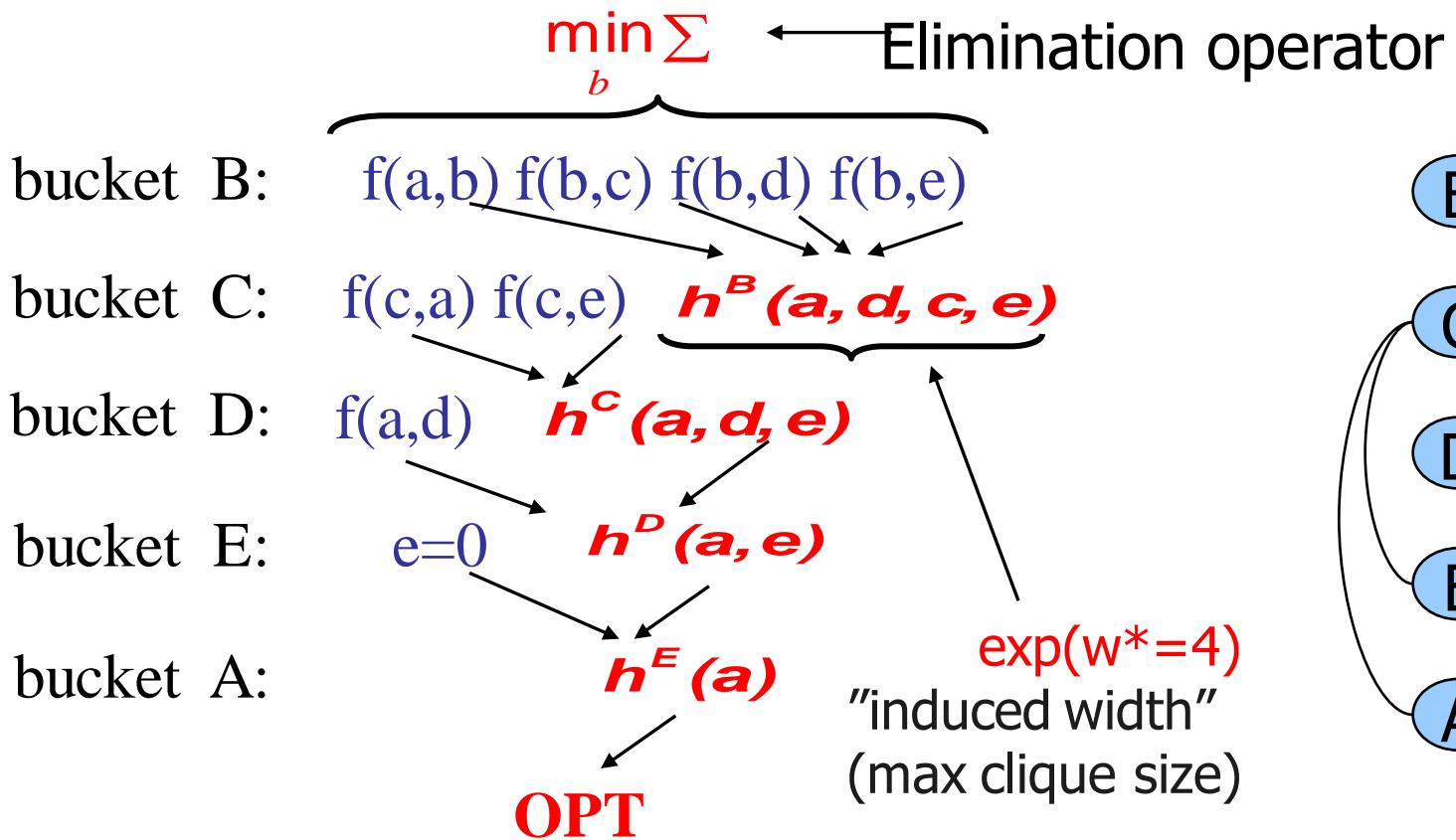
Return (a', b', c', d', e')

Complexity

Algorithm **elim-opt** (Dechter, 1996)

Non-serial Dynamic Programming (Bertele & Brioschi, 1973)

$$OPT = \min_{a,e,d,c,b} F(a,b) + F(a,c) + F(a,d) + F(b,c) + F(b,d) + F(b,e) + F(c,e)$$



Complexity of Bucket Elimination

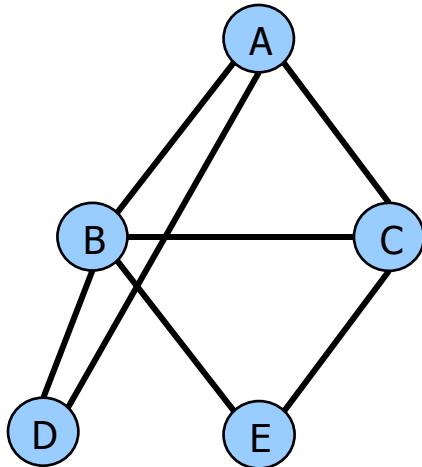
Bucket-Elimination is **time** and **space**

r = number of functions

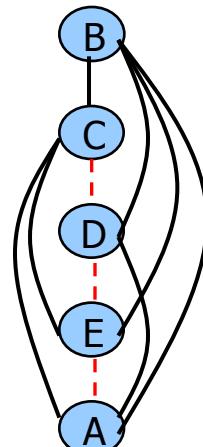
$$O(r \exp(w^*(d)))$$

$w^*(d)$ – the induced width of the primal graph along ordering d

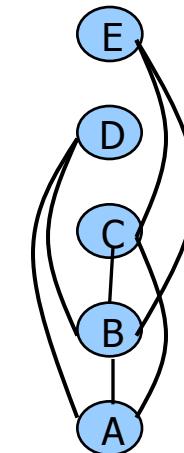
The effect of the ordering:



constraint graph



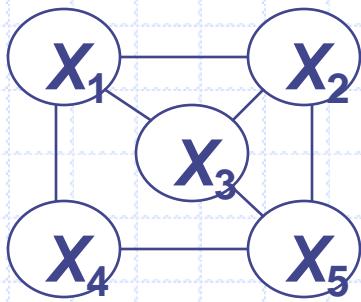
$$w^*(d_1) = 4$$



$$w^*(d_2) = 2$$

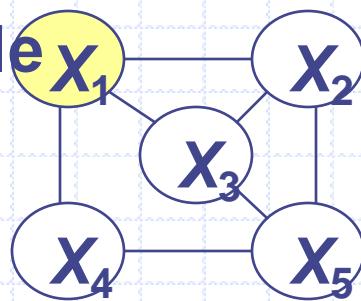
Finding smallest induced-width is hard!

Search Basic Step: Conditioning

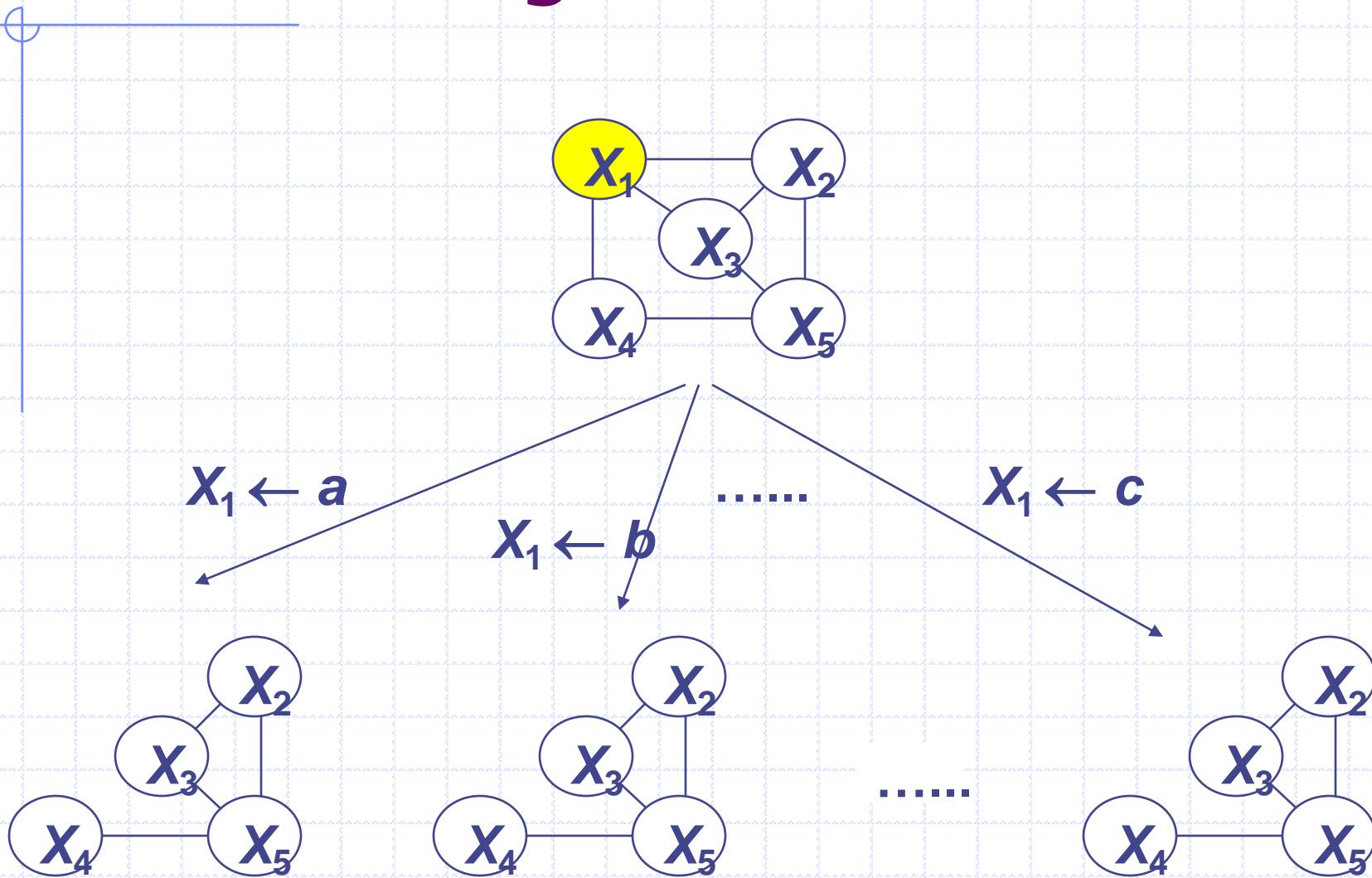


Search Basic Step: Conditioning

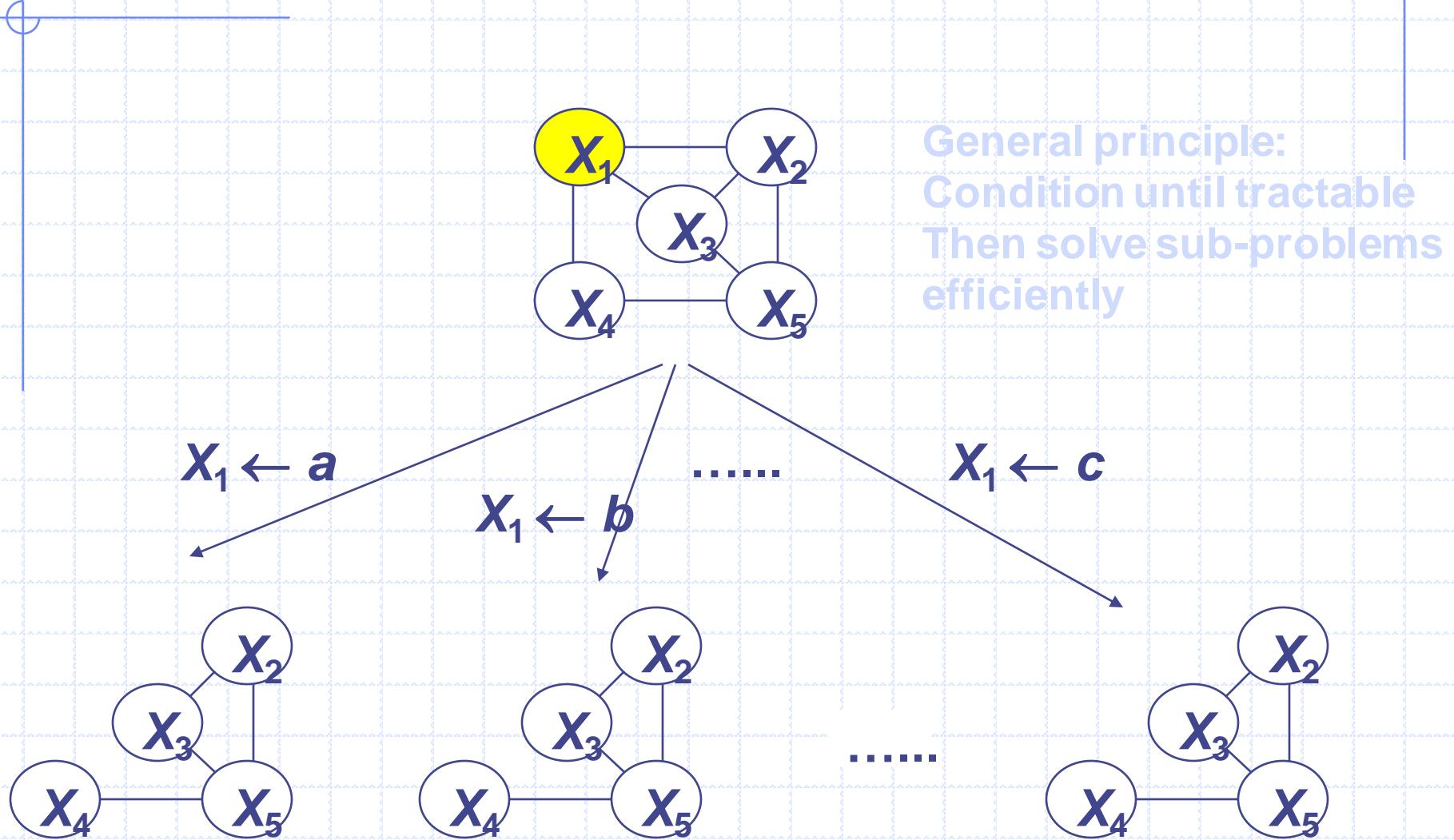
- Select a variable



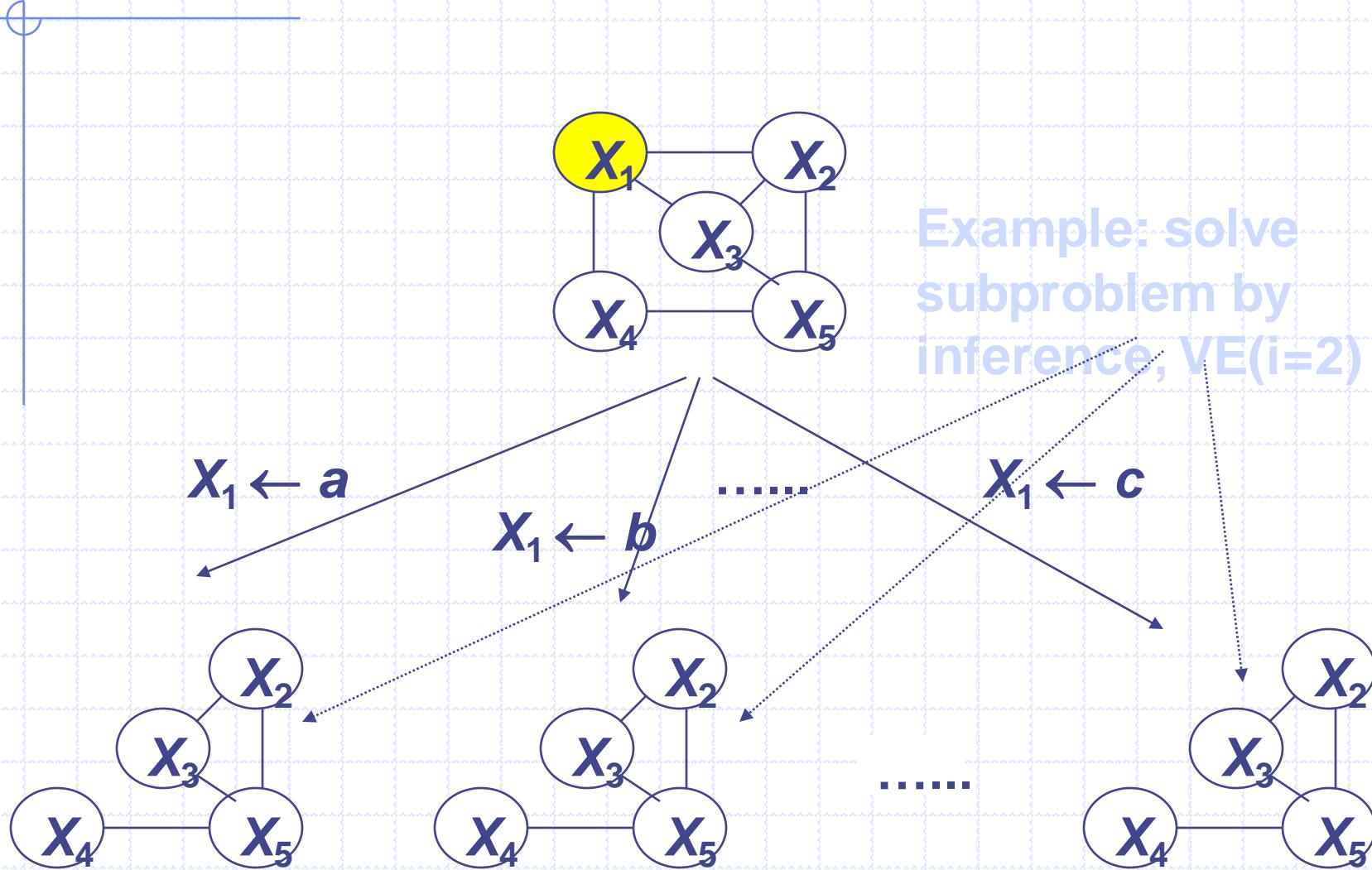
Search Basic Step: Conditioning



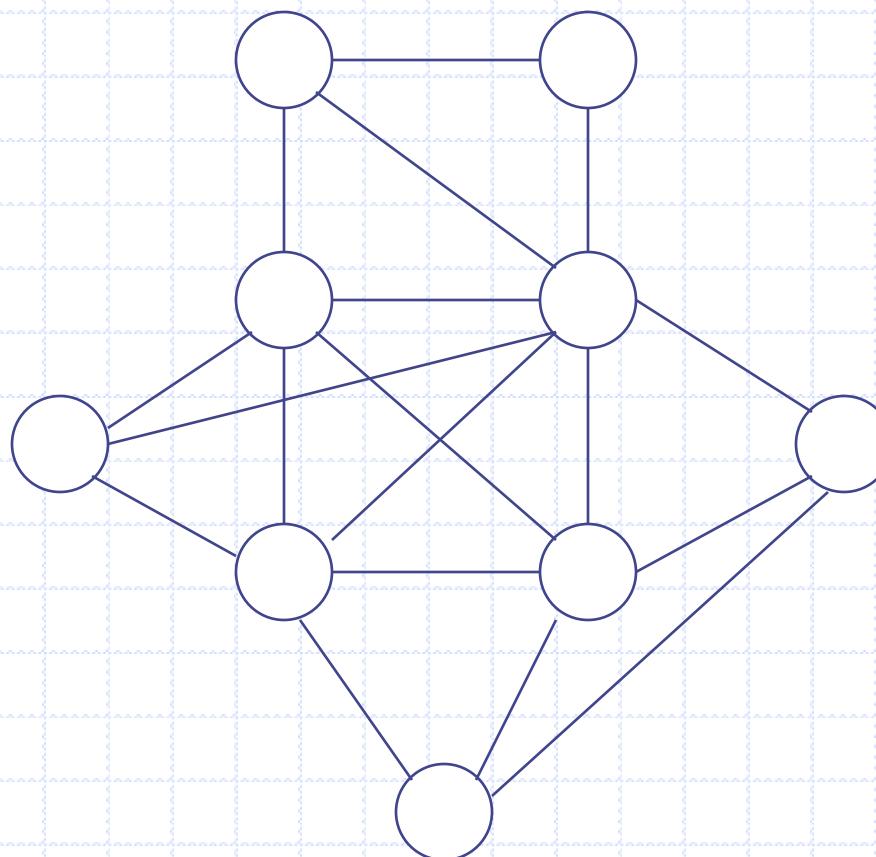
Search Basic Step: Variable Branching by Conditioning



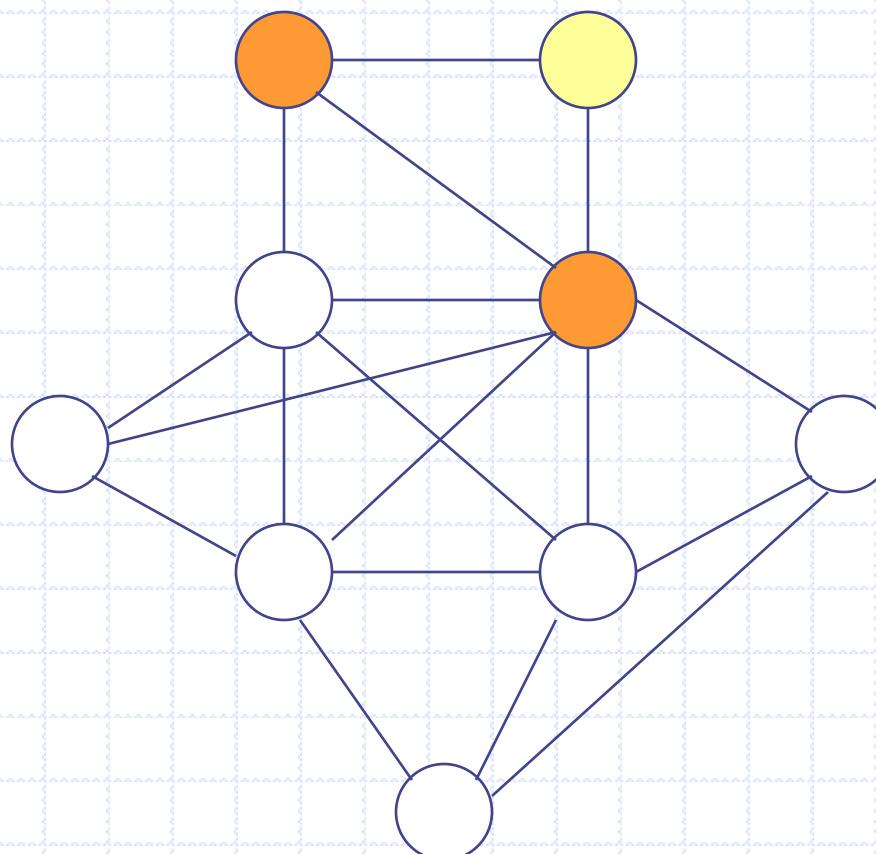
Search Basic Step: Variable Branching by Conditioning



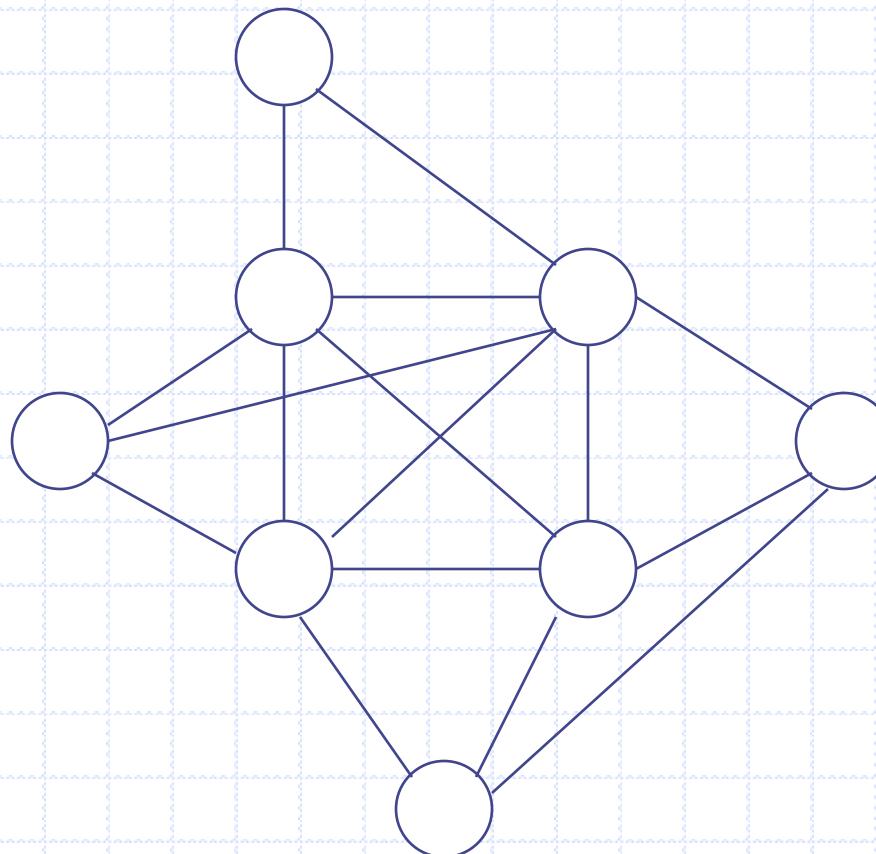
Eliminate First



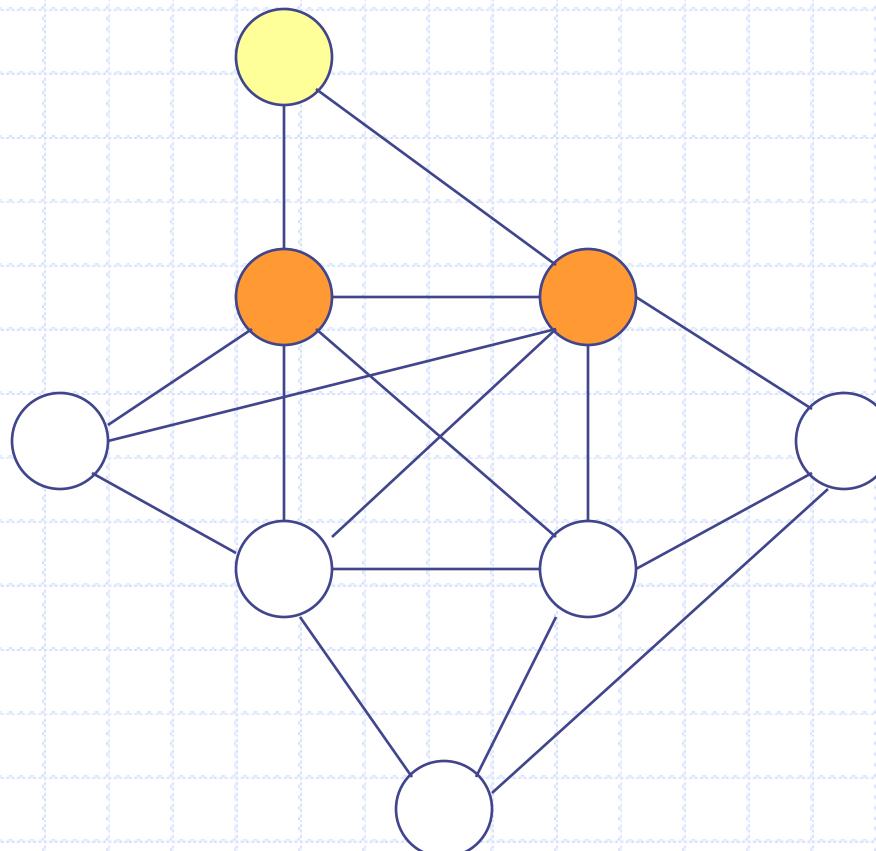
Eliminate First



Eliminate First

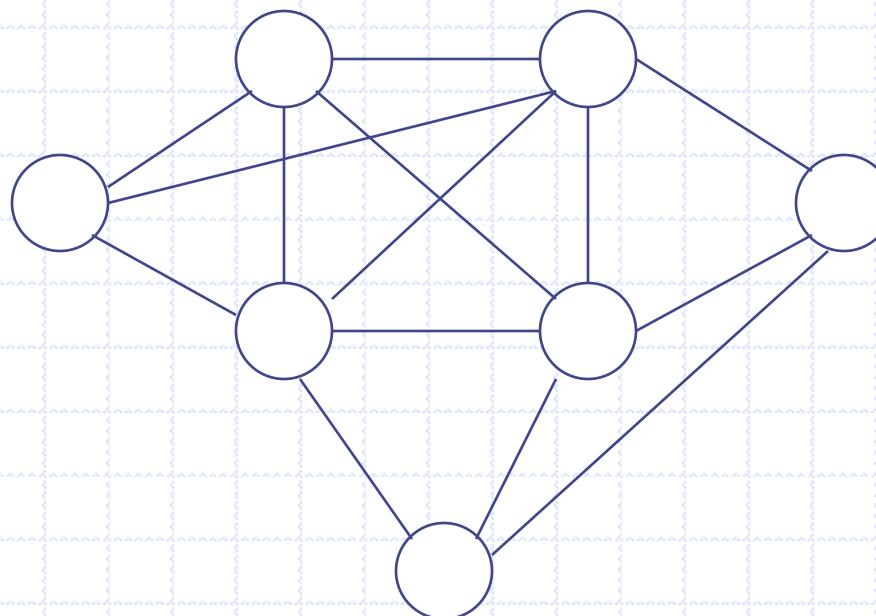


Interleaving Conditioning and Elimination



Interleaving Conditioning and Elimination

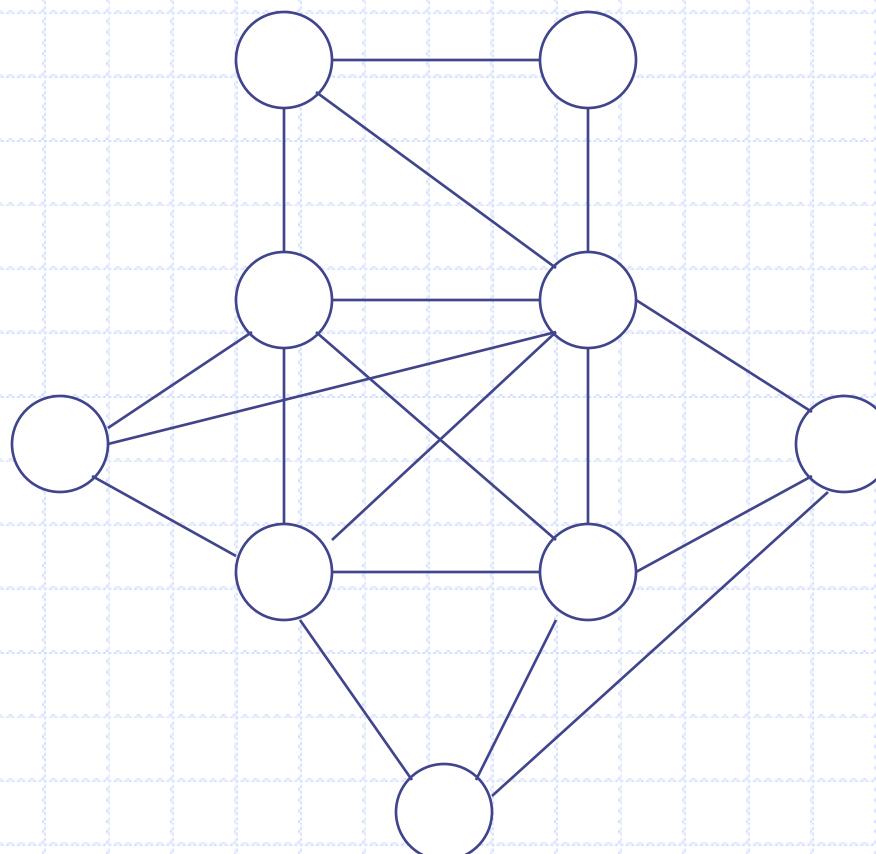
Solve the rest of the problem
by any means



First hybrids: Search & Variable Elimination

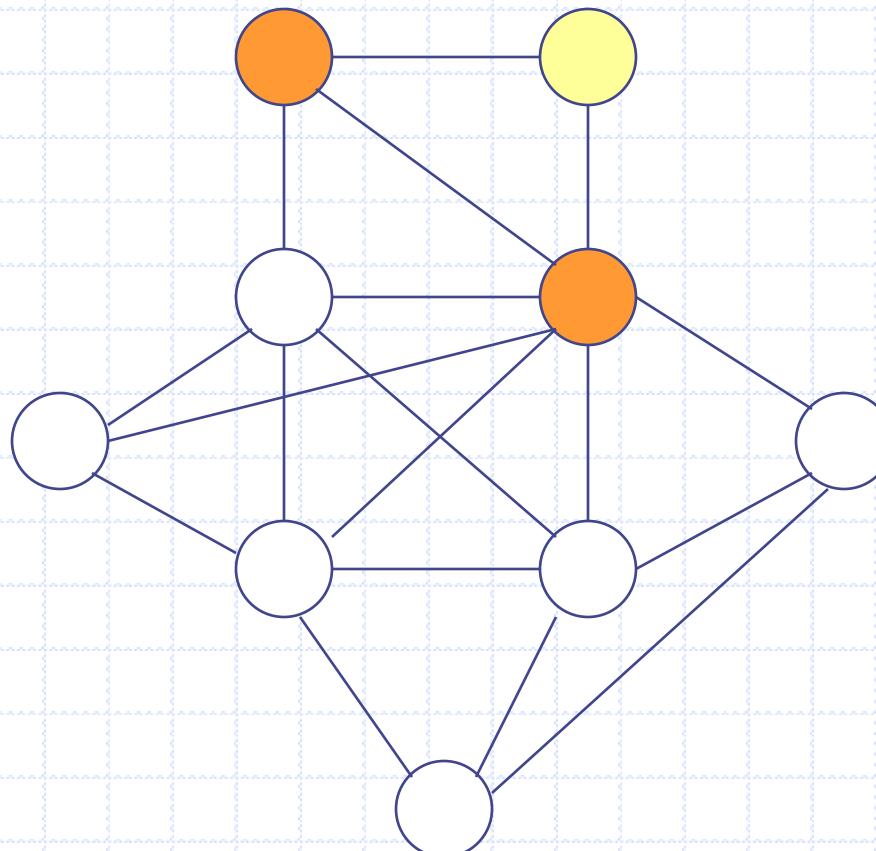
- ◆ Condition, condition, condition ... and then only eliminate (*Cycle-Cutset*)
- ◆ Eliminate, eliminate, eliminate ... and then only search
- ◆ Interleave conditioning and elimination

Interleaving Conditioning and Elimination BB-VE(2) (Larrosa & Dechter, CP 2002)



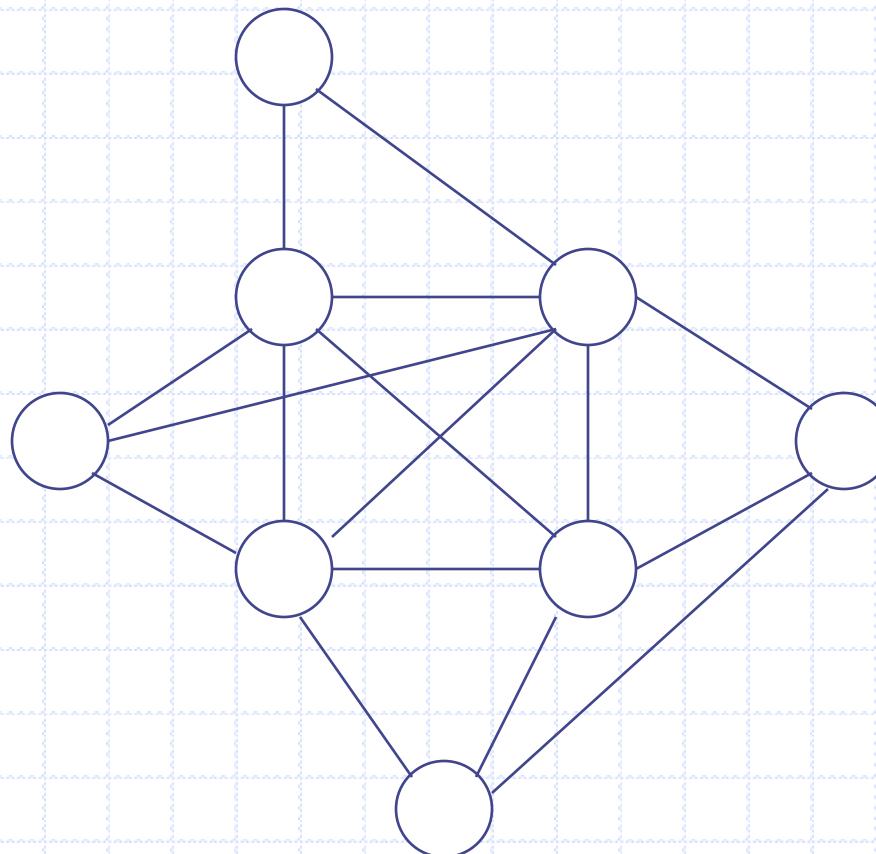
Interleaving Conditioning and Elimination

BB-VE(2)



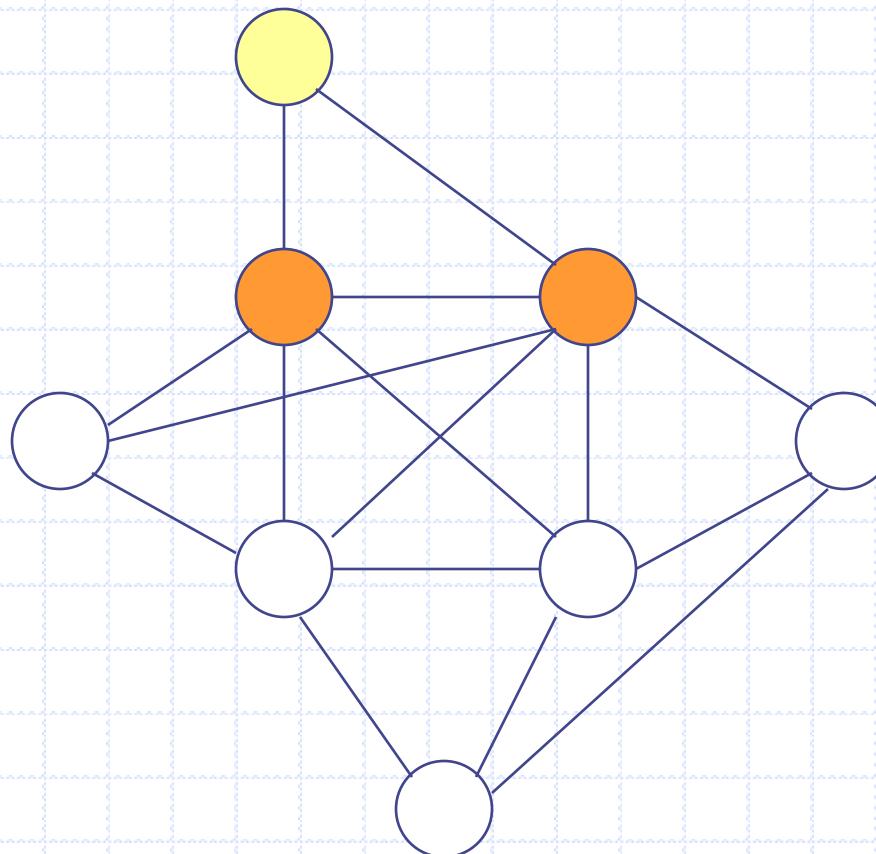
Interleaving Conditioning and Elimination

BB-VE(2)



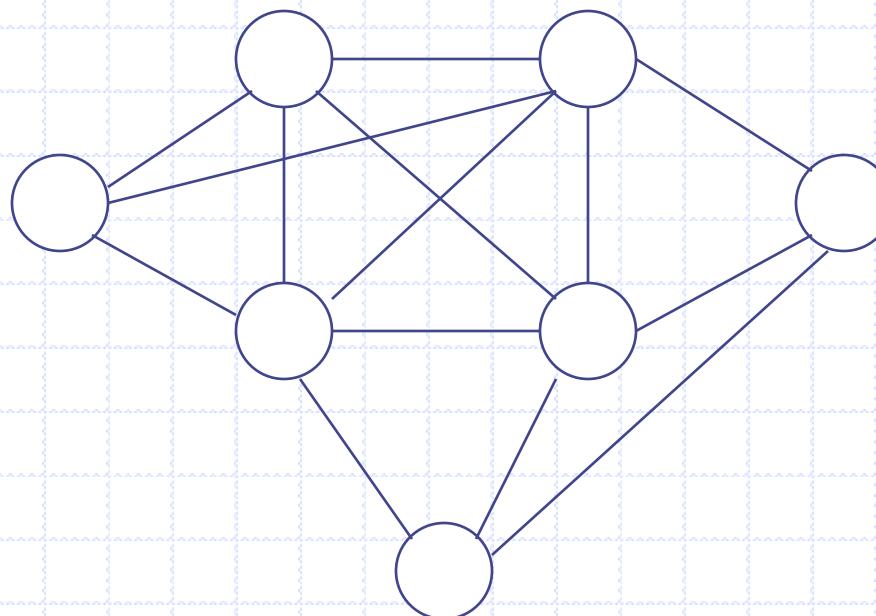
Interleaving Conditioning and Elimination

BB-VE(2)



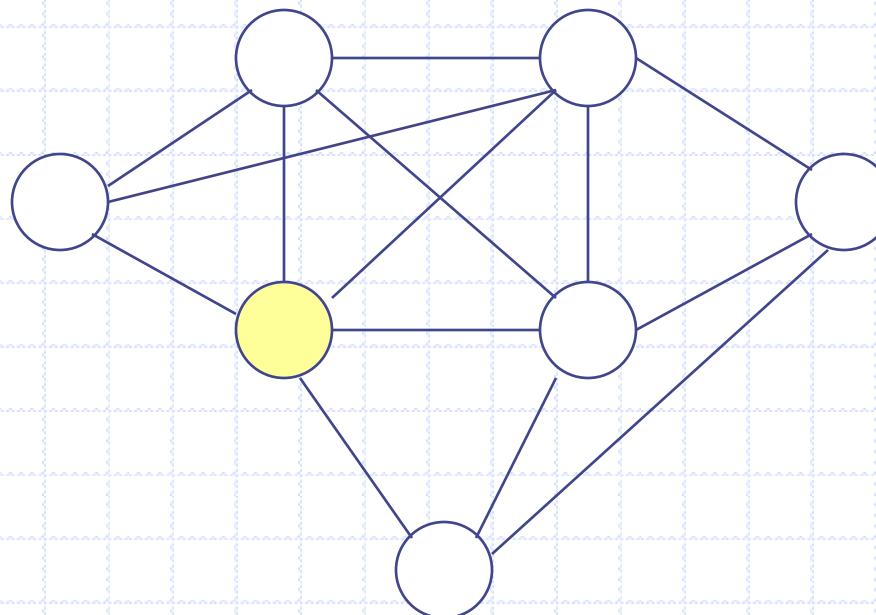
Interleaving Conditioning and Elimination

BB-VE(2)



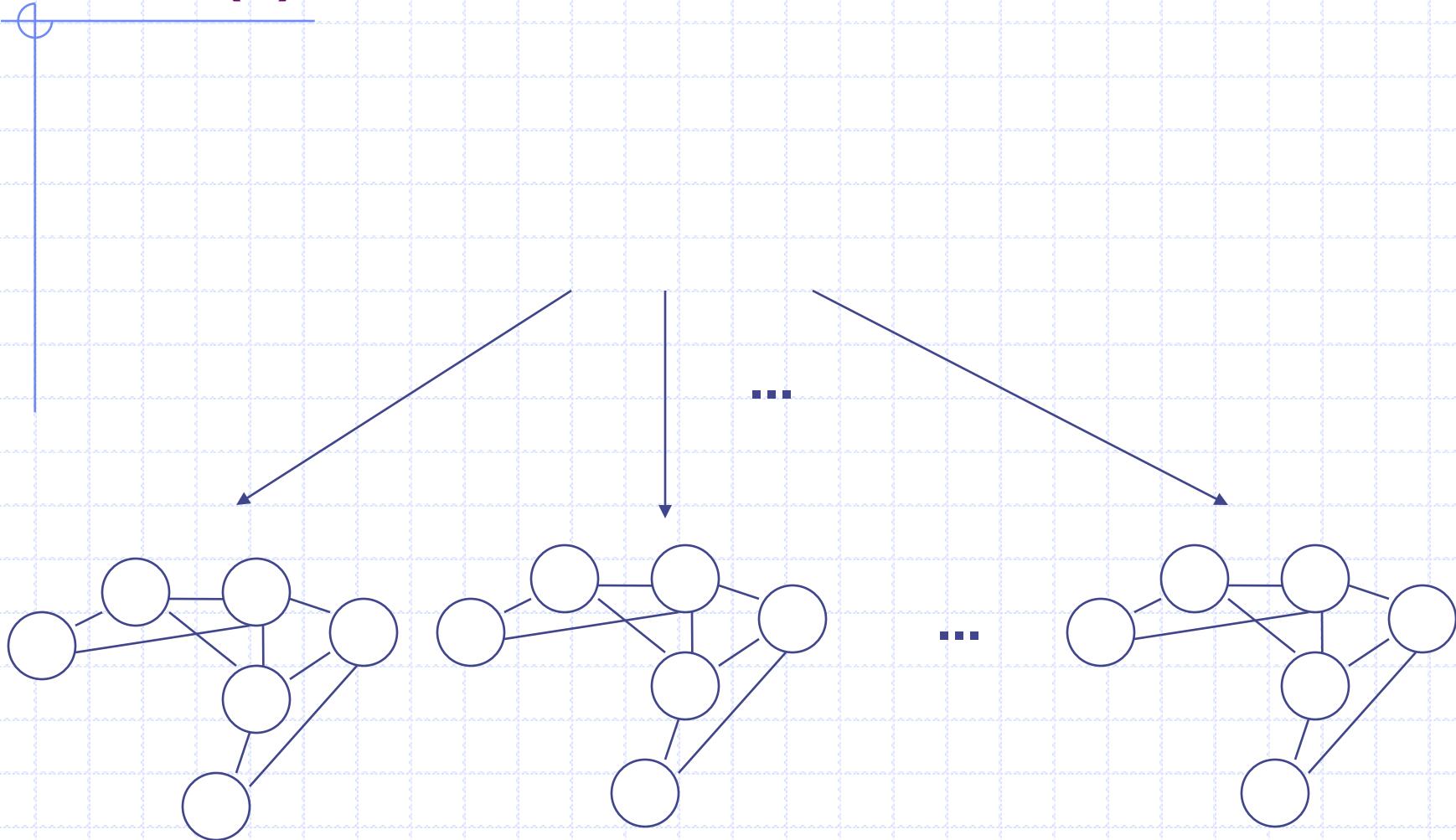
Interleaving Conditioning and Elimination

BB-VE(2)

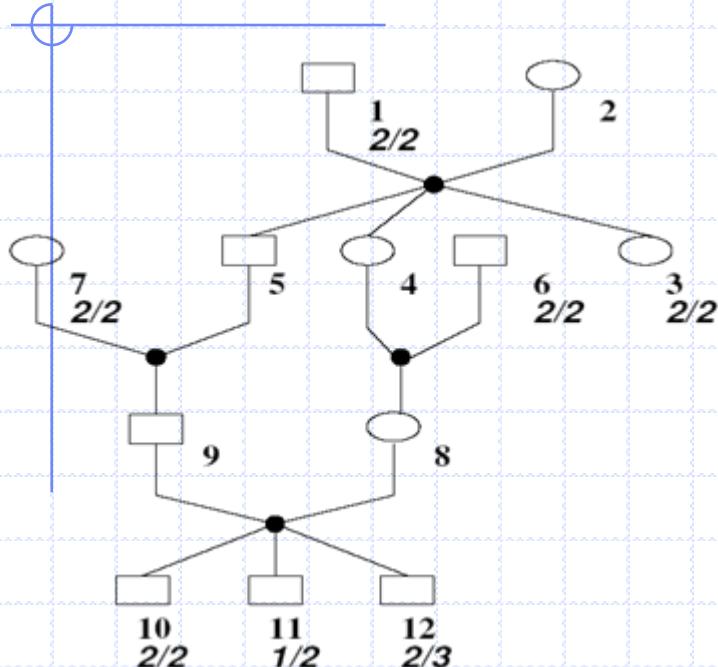


Interleaving Conditioning and Elimination

BB-VE(2)



Mendelian error detection



- Given a pedigree and partial observations (genotypings)
- Find the **erroneous genotypings**, such that their removal restores consistency

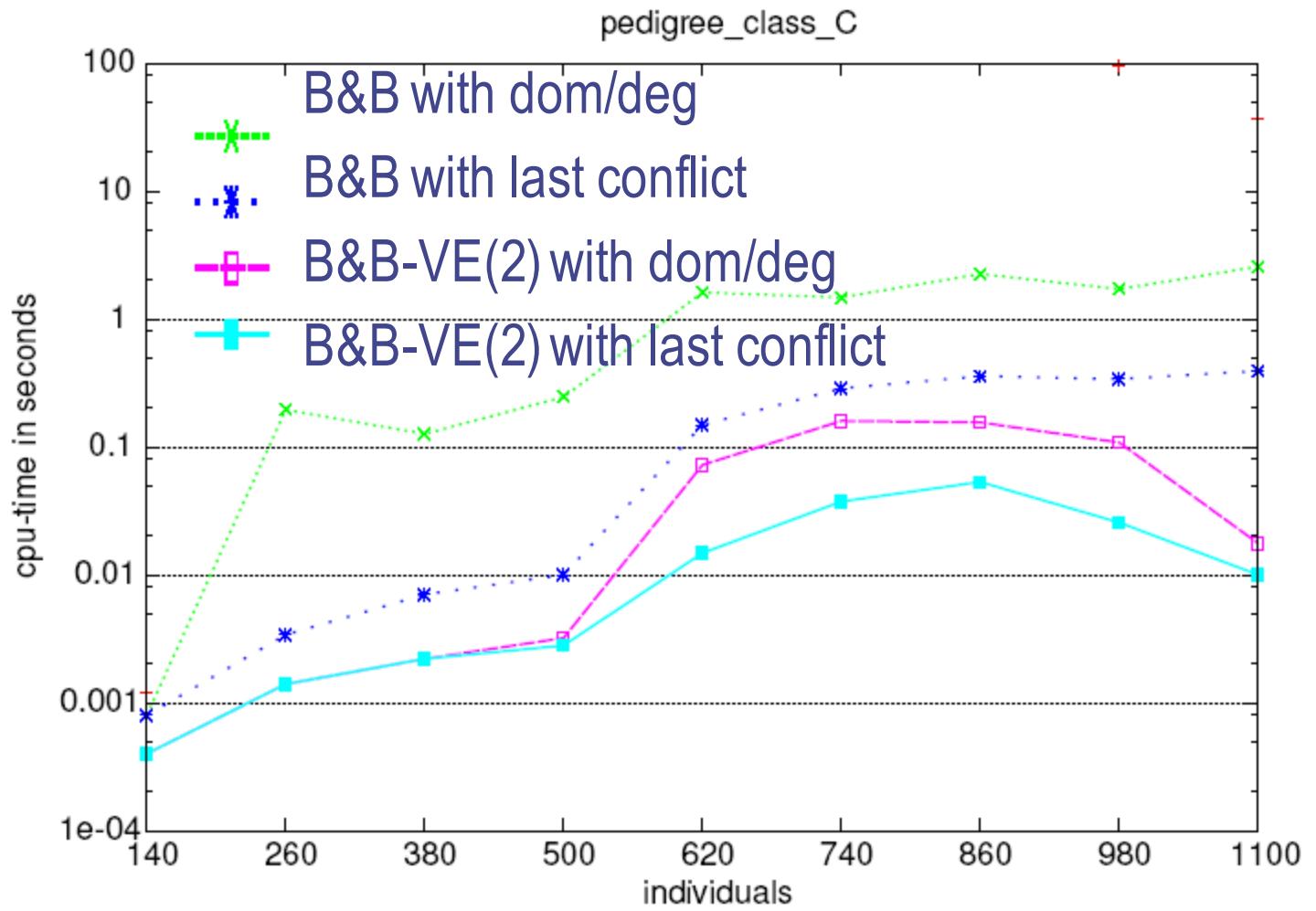
- Checking consistency is NP-complete (Aceto et al., *Comp. Sci. Tech.* 2004)
- Minimize the number of genotypings to be removed
- Maximize the joint probability of the true genotypes (MPE)

Pedigree problem size:

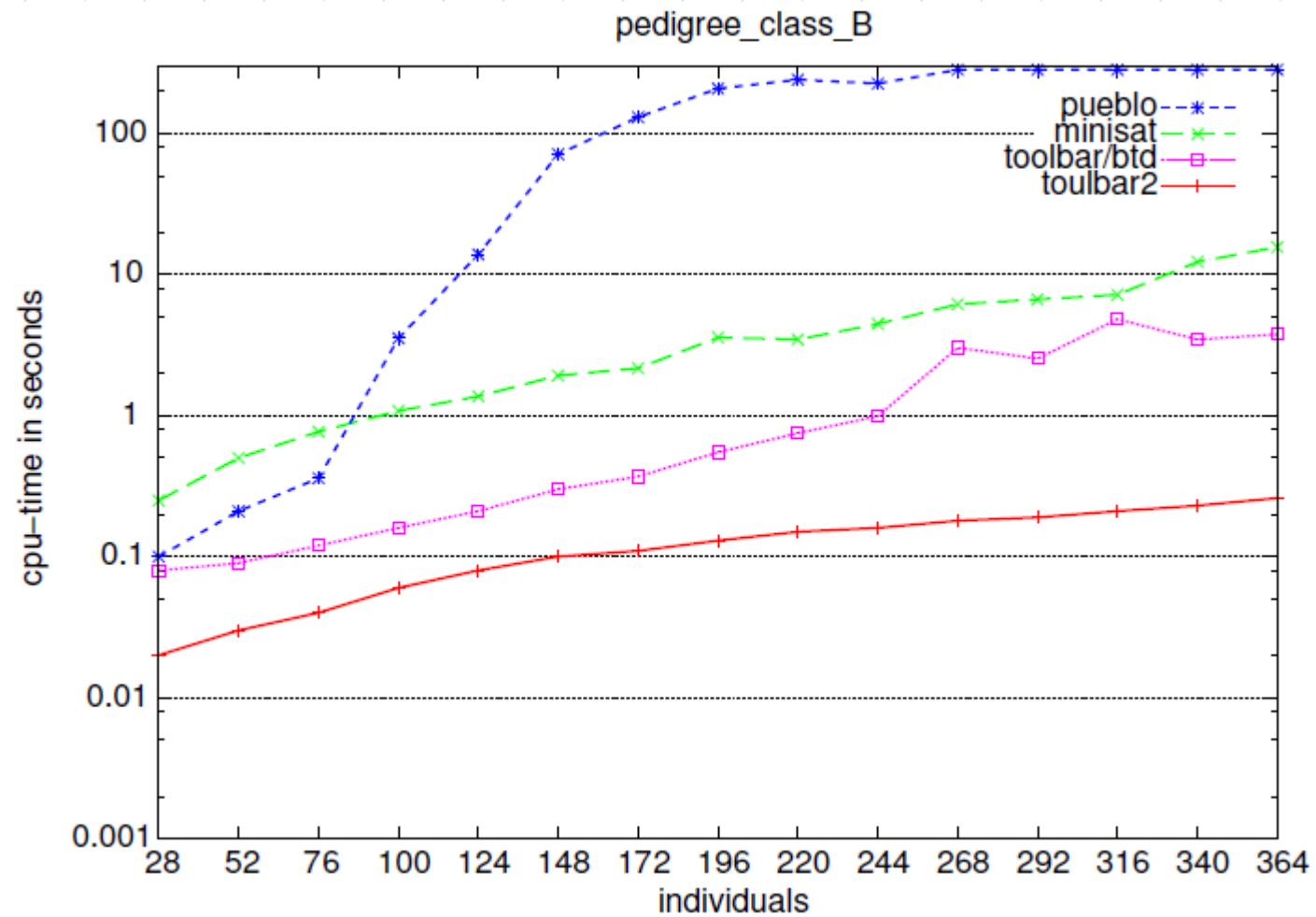
$$n \leq 20,000 ; d = 3-66 ; e(3) \leq 30,000$$

Pedigree

- toulbar2 v0.5 with EDAC3 and binary branching
- Minimize the number of genotypings to be removed
- CPU time in seconds to find and prove optimality on a linux PC 3 GHz with 16 GB



Pedigree



Pedigree

- toulbar2 with last conflict
- Minimize the number of genotypings to be removed
- CPU time in seconds to find and prove optimality

B&B-VE(2)

	ind	vars	genotyped	alleles	nf	ngen	treewidth ub		errors	time	nodes
<i>eye</i>	36	36	28	6	11	4	2		1	0.02	0
<i>cancer</i>	49	48	37	8	18	5	2		1	0.21	0
<i>parkinson</i>	37	34	13	4	7	7	5		0	0	6
<i>berrichon_{1nc}</i>	129516	9947	2448	4	8821	17	262		2	4.73	8805
<i>berrichon₁</i>	129516	10017	2483	4	8786	17	330		23	5.81	8384
<i>berrichon_{2nc}</i>	27255	19337	10215	4	4719	19	-		41	5.89	6170
<i>berrichon₂</i>	27255	19562	10215	4	2381	19	-		106	17.23	15445
<i>langlade₁</i>	1355	1209	711	9	298	13	84		38	12.28	391
<i>langlade₂</i>	1355	1223	715	7	298	13	82		89	60.56	17857
<i>langlade₃</i>	1355	1258	787	5	298	13	85		39	14.19	6731
<i>langlade₄</i>	1355	1186	672	8	298	13	83		43	59.7	3520
<i>moissac₁</i>	283	260	183	2	81	5	6		0	0	5
<i>moissac₂</i>	283	244	167	7	81	5	6		0	0.51	6
<i>moissac₃</i>	283	225	151	3	81	5	6		0	0	4
<i>moissac₄</i>	283	256	179	2	81	5	6		0	0	5
<i>moissac₅</i>	283	237	161	8	81	5	6		0	1.02	5
<i>moissac₆</i>	283	201	131	11	81	5	5		0	5.64	6

Solving methods

BTD, AND/OR graph search

Time: $\exp(\text{treewidth})$

Space: $\exp(\text{treewidth})$

Search: Conditioning

Time: $\exp(n)$

Space: linear

BB-VE

Time: $\exp(n)$

Space: linear

Inference: Elimination

Time: $\exp(\text{treewidth})$

Space: $\exp(\text{treewidth})$

Complete

Depth-First
Branch & Bound

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Mini-bucket(i)

Hybrids

Complete

Variable Elimination

Cluster Tree Elimination

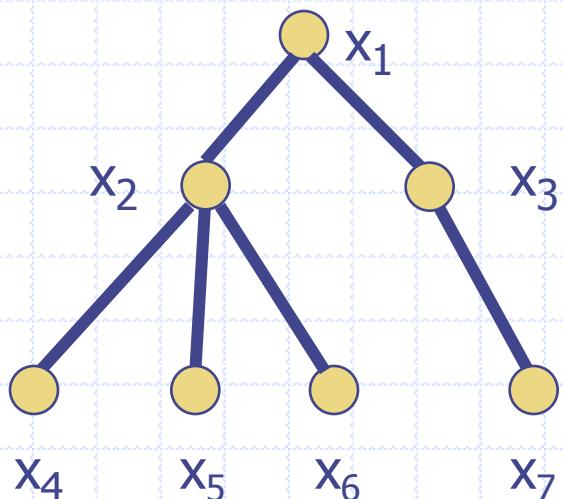
Second hybrids: Search & Cluster Tree Elimination

- ◆ Depth-First Branch and Bound
exploiting a tree decomposition with:
 - A restricted variable ordering
 - Graph-based backjumping
 - Graph-based learning

⇒ Lazy elimination of subproblems
using search

Structural tractability

Tree-structured binary WCSPs are tractable



Time complexity $O(e d^2)$
Space complexity $O(n d)$

*n: number of variables
d: maximum domain size
e: number of cost functions*

Proceed from the leaf nodes to a chosen root node

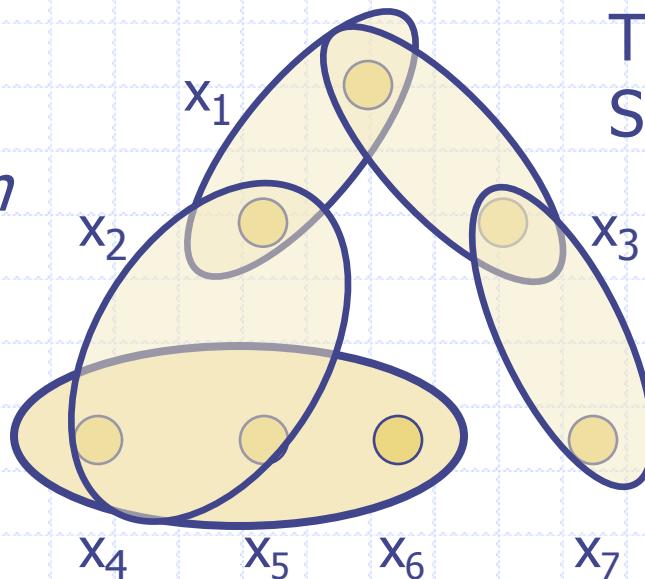
Project out leaf nodes by minimising over possible assignments

e.g. add $\text{PROJ}_{x_2}(f_{x_2} + f_{x_2, x_4})$ to f_{x_2}

Structural tractability

- ◆ Acyclic r-ary WCSPs are tractable

*Each cost function
is represented by
an hyperedge*



Time complexity in $O(e^{d^r})$
Space complexity in $O(n^{d^s})$

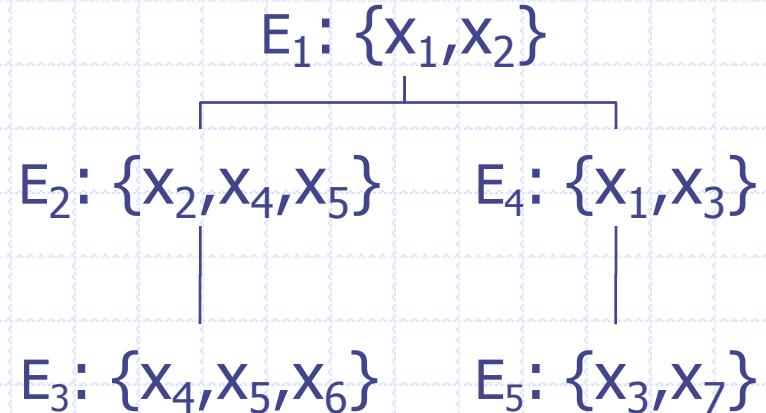
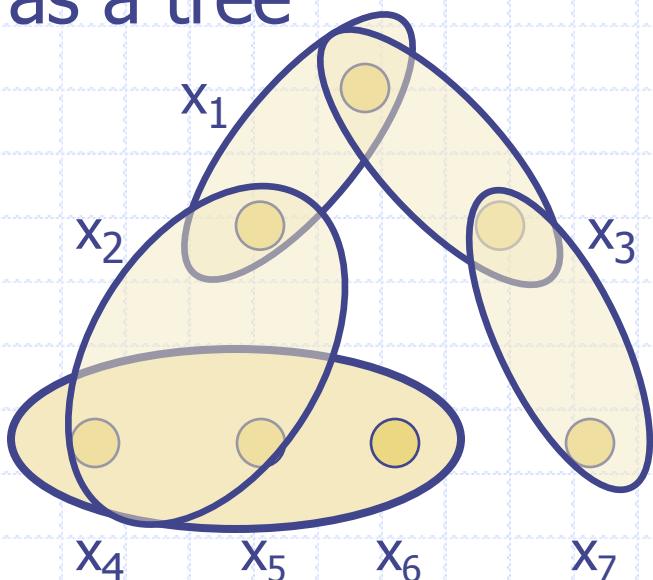
*r: bounded max arity
s: $\max\{|U \cap V| : f_U \neq f_V\}$*

Proceed from the *leaf* hyperedges to a chosen root hyperedge

e.g. add $\text{PROJ}_{x_4,x_5}(f_{x_6} + f_{x_4,x_5,x_6})$ to f_{x_4,x_5}

Tree decomposition

- ◆ A collection of subsets of X (clusters) organized as a tree

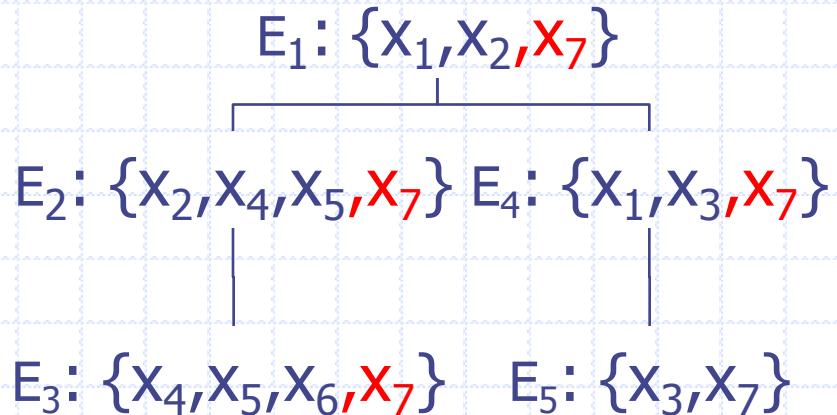
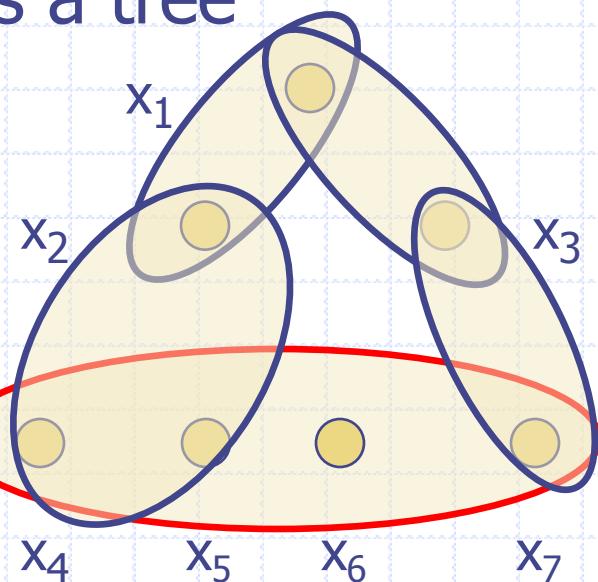


Properties:

- $\bigcup E_i = X$
- $\forall f_S \in F, \exists E_i$ such that $S \in E_i$
- $\forall x \in X, \{E_i : x \in E_i\}$ forms a connected subtree

Tree decomposition

- ◆ A collection of subsets of X (clusters) organized as a tree

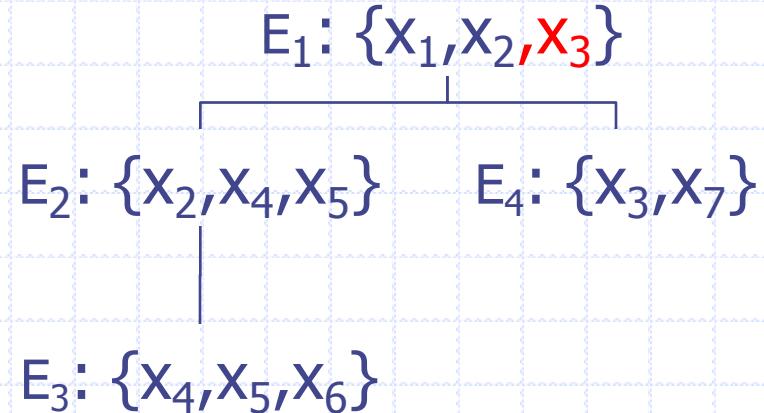
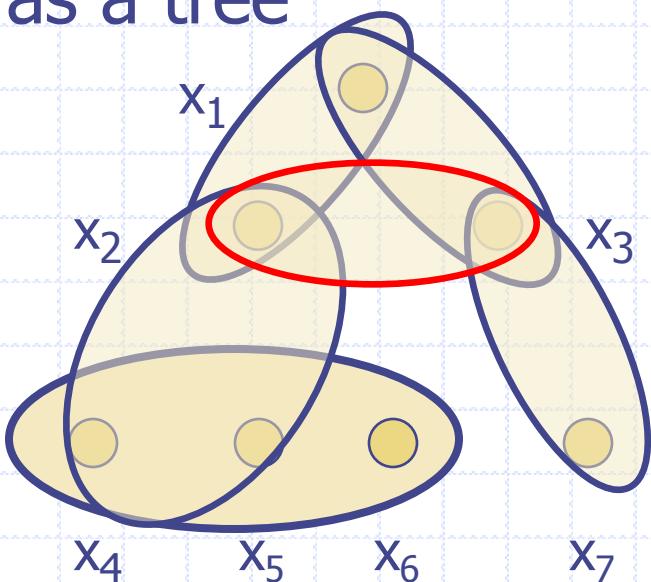


Properties:

- $\cup E_i = X$
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- $\forall x \in X, \{E_i : x \in E_i\}$ forms a **connected subtree**

Tree decomposition

- ◆ A collection of subsets of X (clusters) organized as a tree

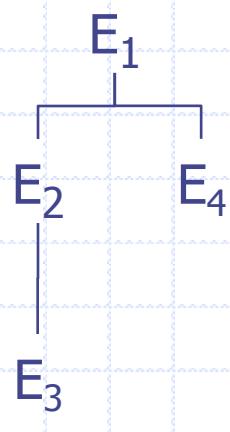
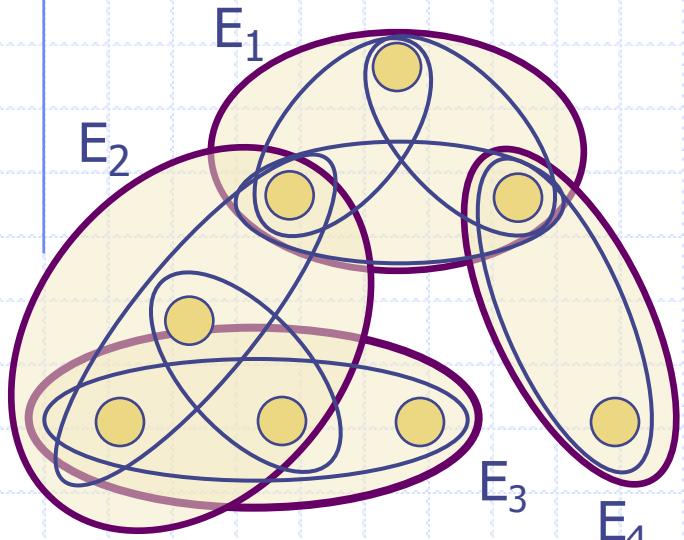


Properties:

- $\bigcup E_i = X$
- $\forall c_S \in P, \exists E_i$ such that $S \in E_i$
- $\forall x \in X, \{E_i : x \in E_i\}$ forms a connected subtree

Tree decomposition

Bounded treewidth VCSPs are tractable



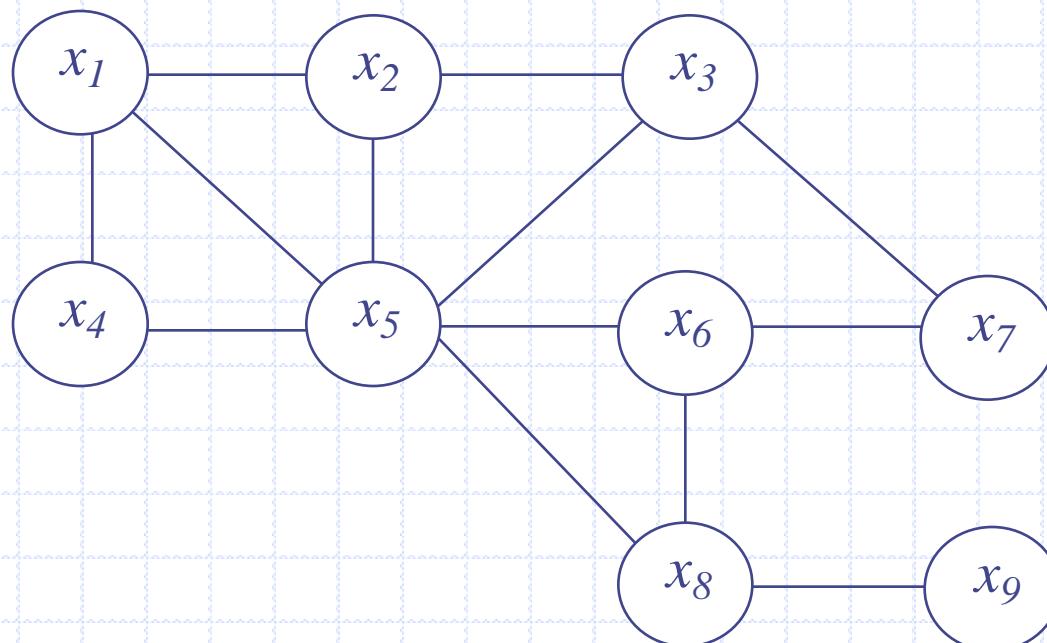
Time complexity $O(e^{d^{w+1}})$
Space complexity $O(n d^s)$

w: bounded treewidth
 $= \max |E_i| - 1$

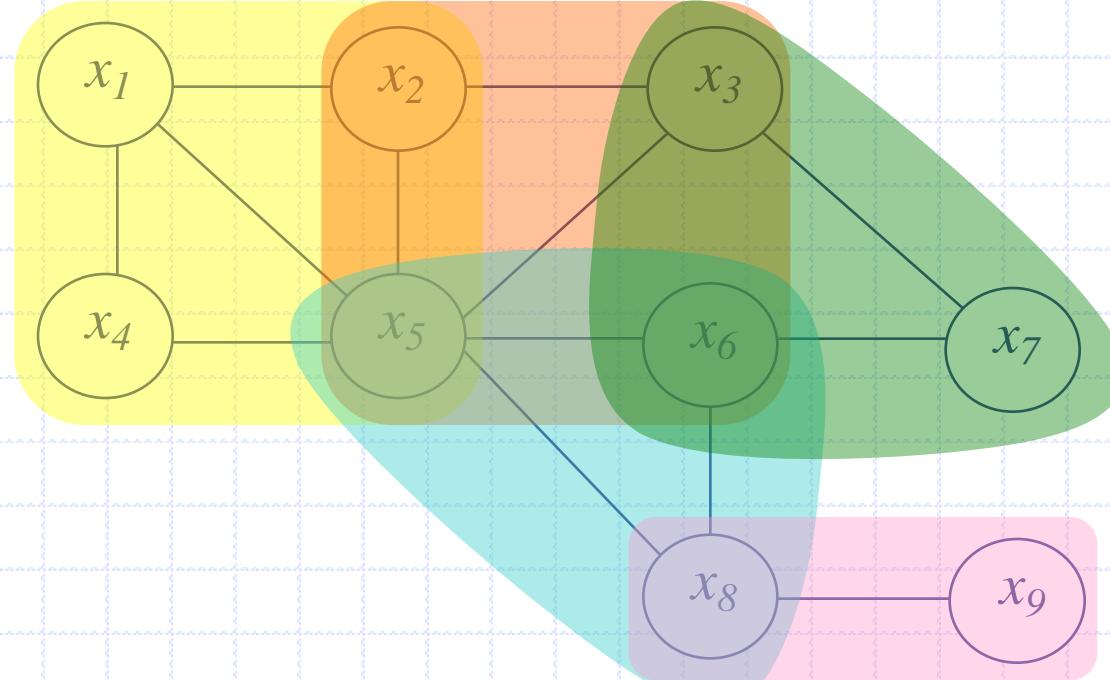
s: $\max \{|E_i \cap E_j| : i \neq j\}$

Finding a tree decomposition with minimum w^* is NP-hard!

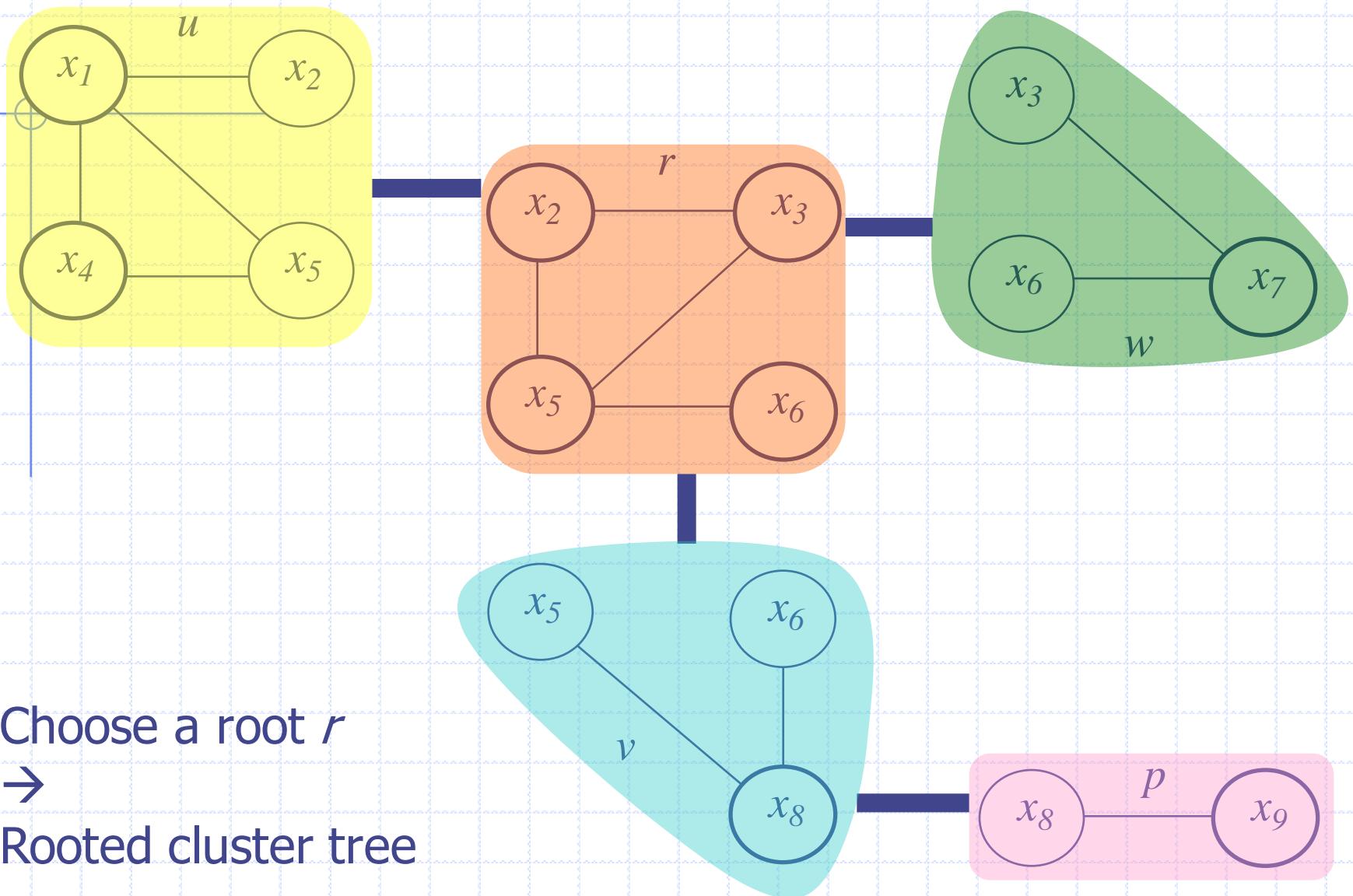
Cluster tree elimination



Cluster tree elimination

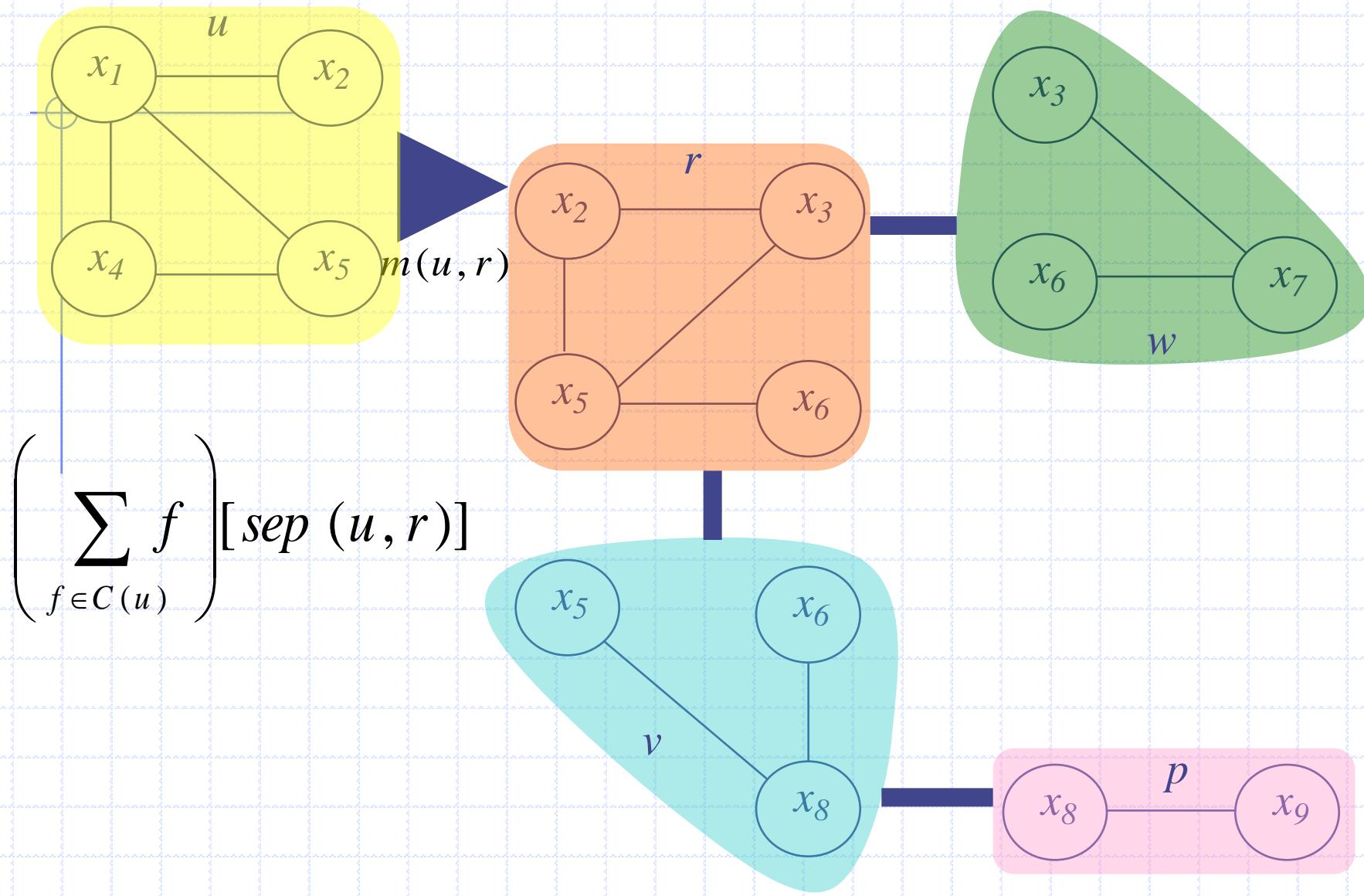


Tree decomposition

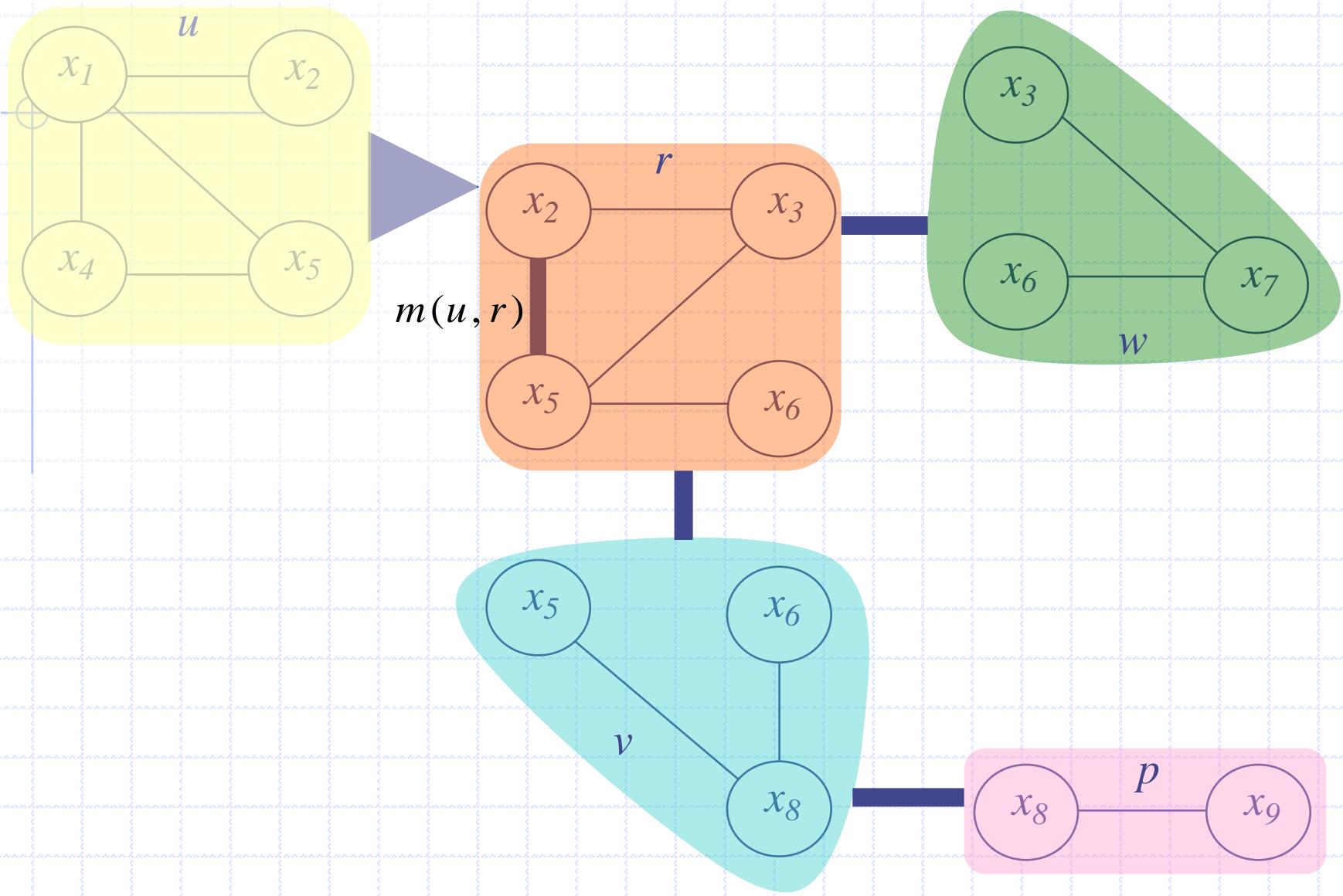


Choose a root r
 \rightarrow
 Rooted cluster tree

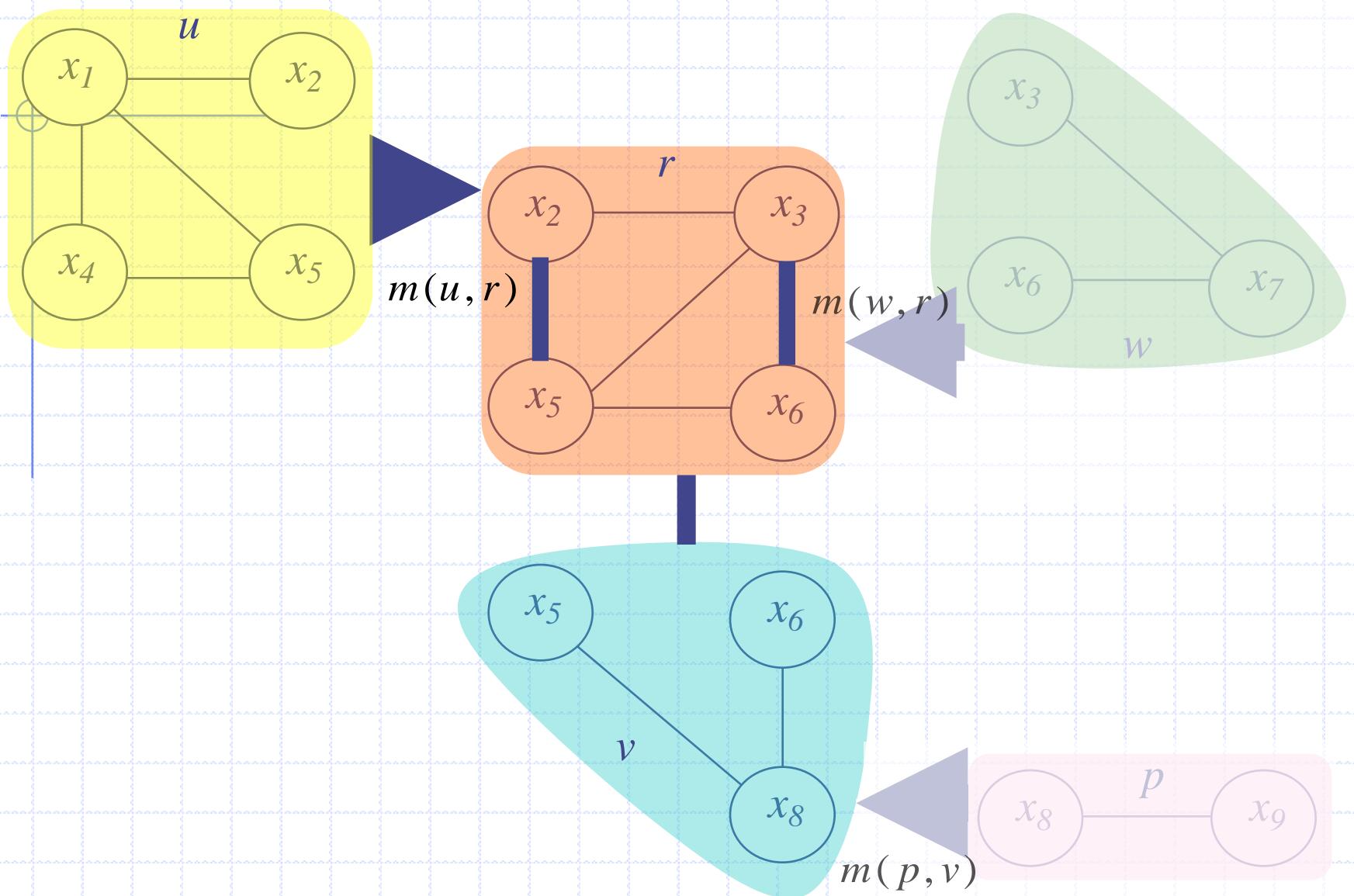
Cluster Tree Elimination



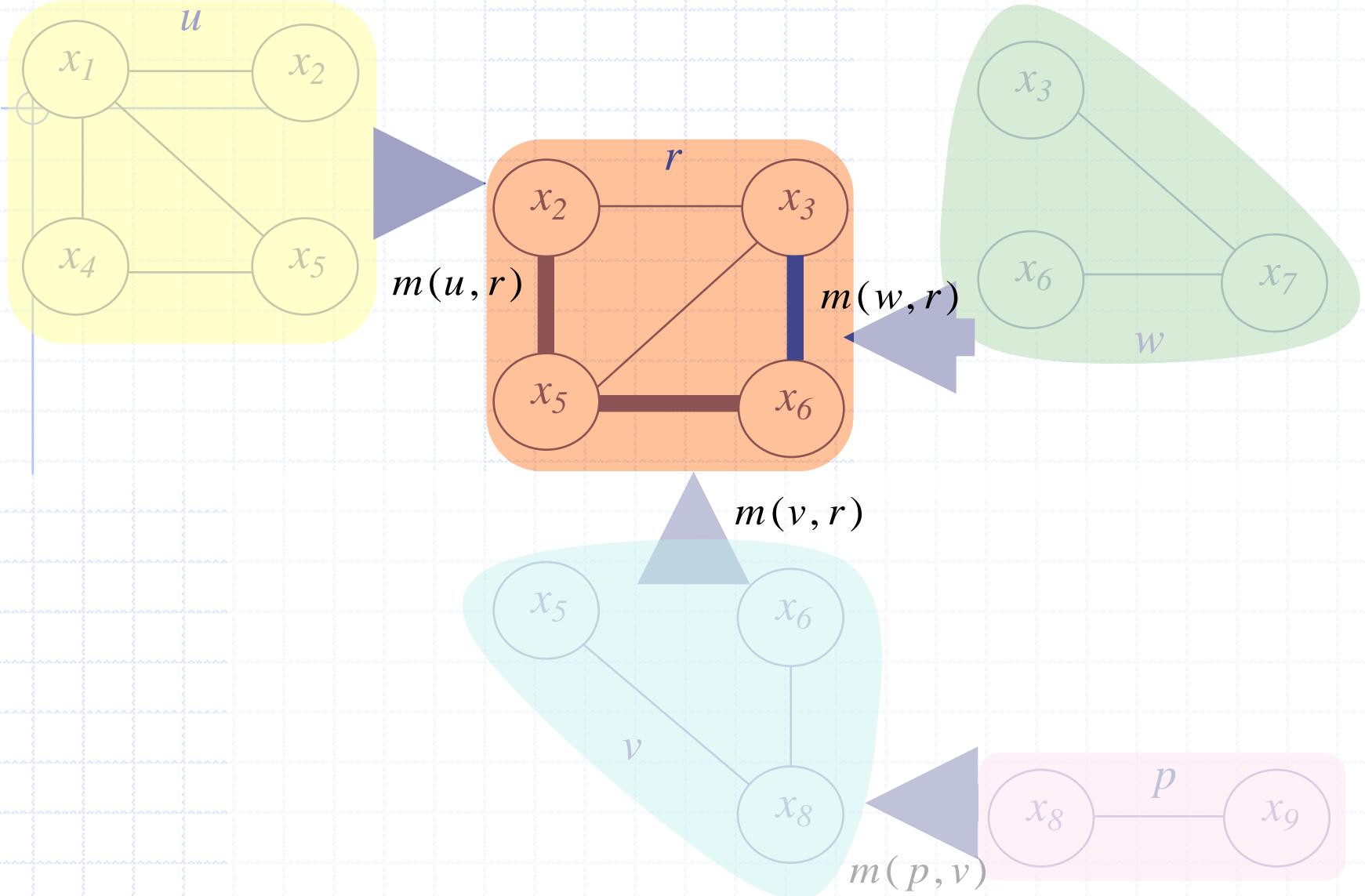
Cluster Tree Elimination



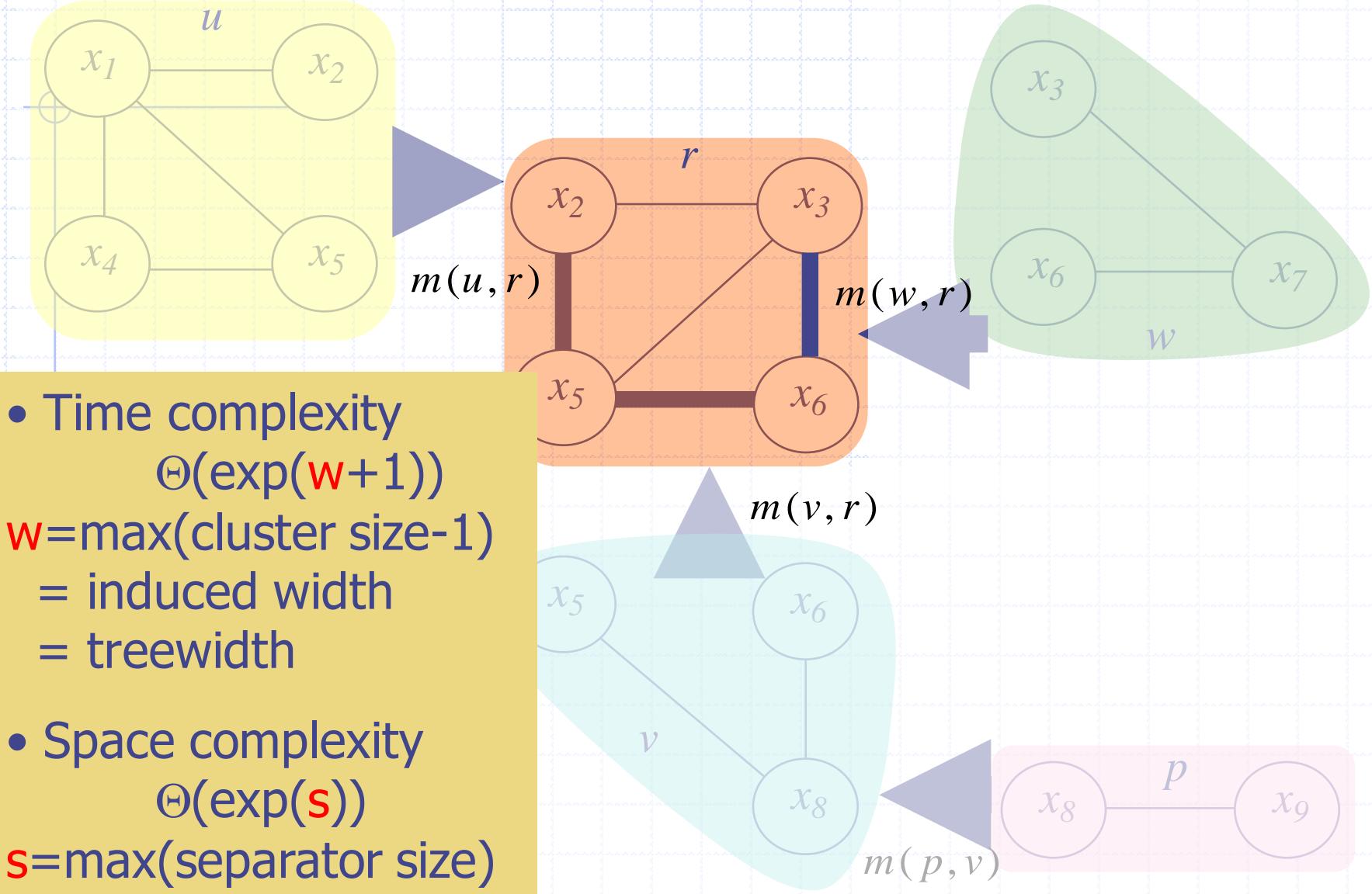
Cluster Tree Elimination



Cluster Tree Elimination



Cluster Tree Elimination



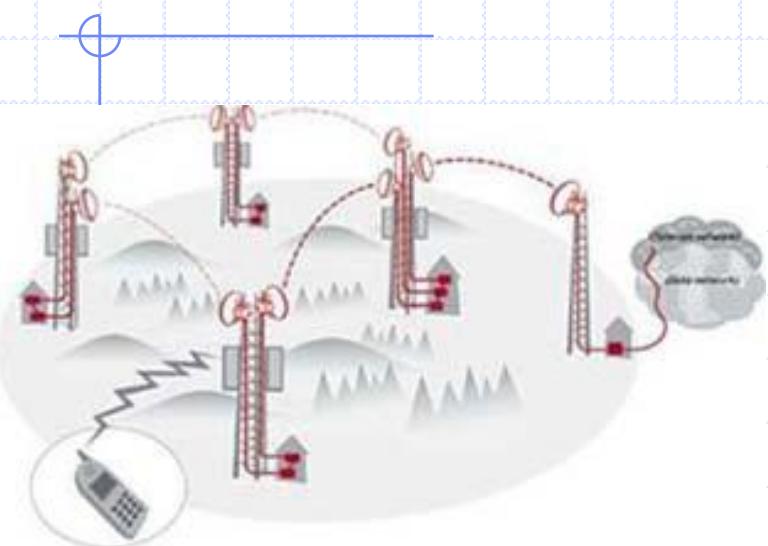
- Time complexity
 $\Theta(\exp(w+1))$
 $w = \max(\text{cluster size}-1)$
= induced width
= treewidth
- Space complexity
 $\Theta(\exp(s))$
 $s = \max(\text{separator size})$
 $\leq w$

History / terminology

- ◆ Operations Research: Non serial **dynamic programming** (Bertelé Brioschi, 72)
- ◆ Databases: Acyclic DB (Beeri et al 1983)
- ◆ Bayesian nets: Join-tree (Pearl 88, Lauritzen et Spiegelhalter 88)
- ◆ Constraint nets: Adaptive Consistency (Dechter and Pearl 88)
- ◆ Genetics: Peeling (Elston and Stewart 71)

Radio Link Frequency Assignment Problem

(Cabon et al., *Constraints* 1999) (Koster et al., *4OR* 2003)



- Given a telecommunication network
- ...find the **best** frequency for each communication link, avoiding interferences

- **Best** can be:
 - Minimize the maximum frequency, no interference (max operator)
 - **Minimize the global interference (sum operator)**
 - Generalizes graph coloring problems: $|f_i - f_j| \geq a$

CELAR problem size: $n=100-458$; $d=44$; $e=1,000-5,000$

Tree decomposition example

CELAR scen06r

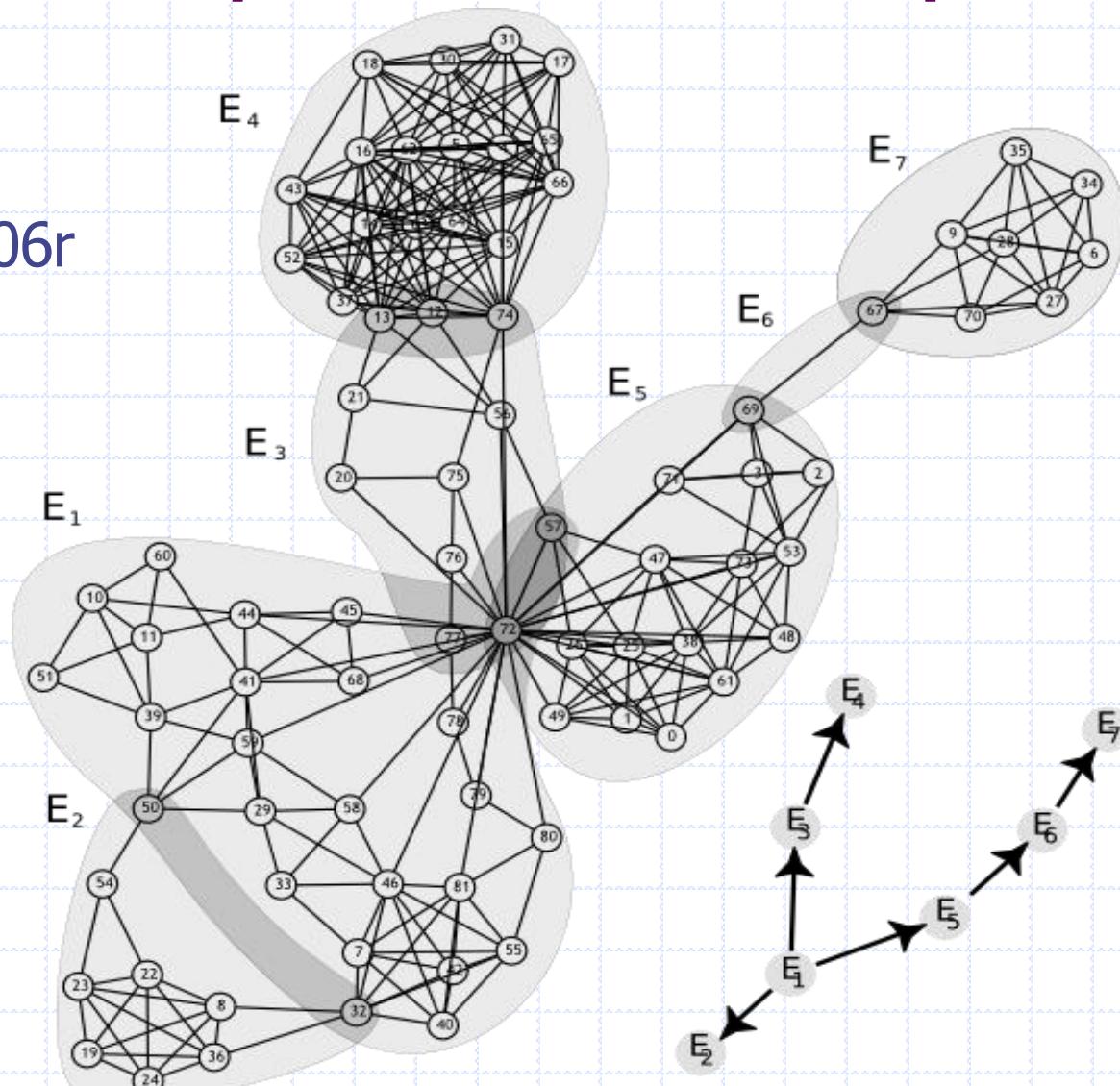
$n = 82$

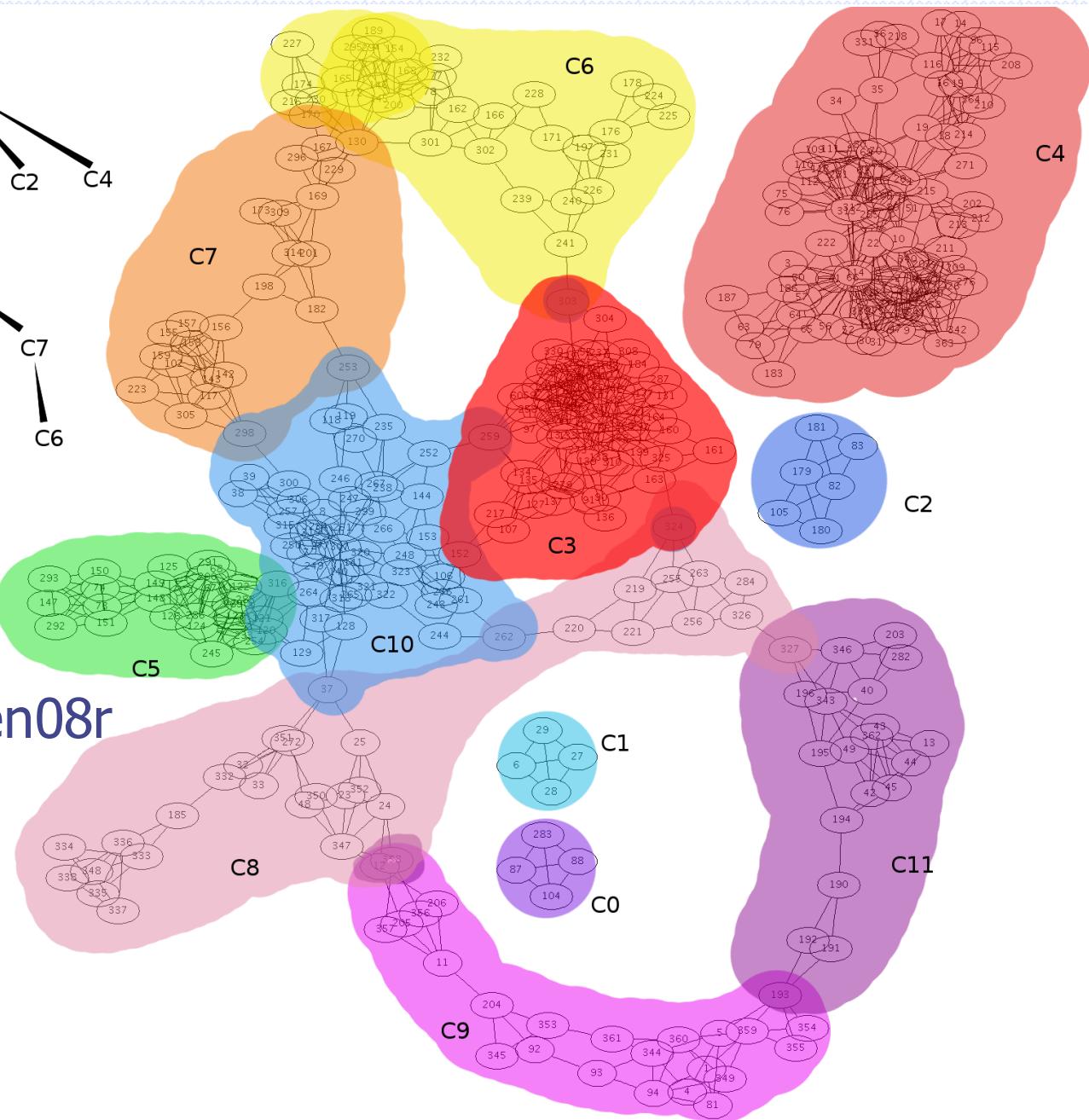
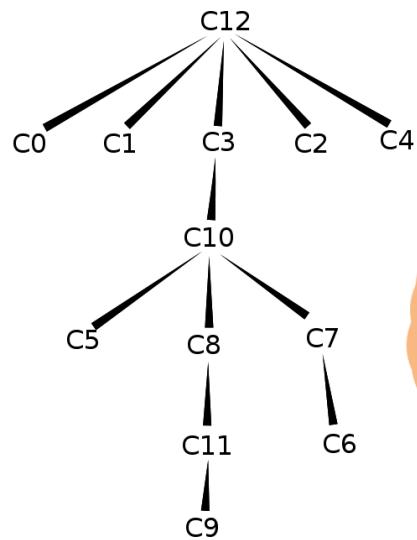
$d = 44$

$e = 327$

$w = 26$

$s = 3$





CELAR scen08r

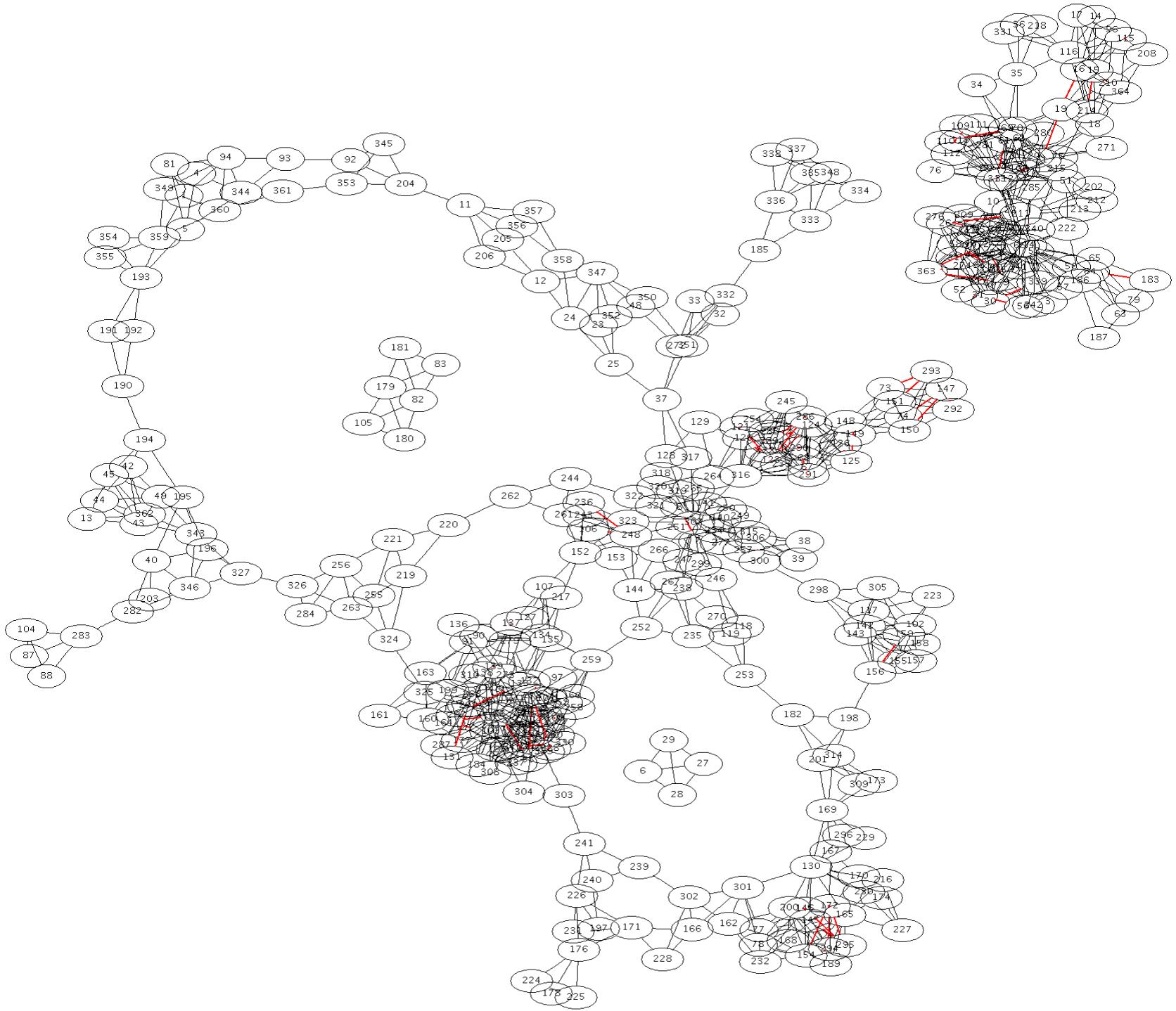
$n = 365$

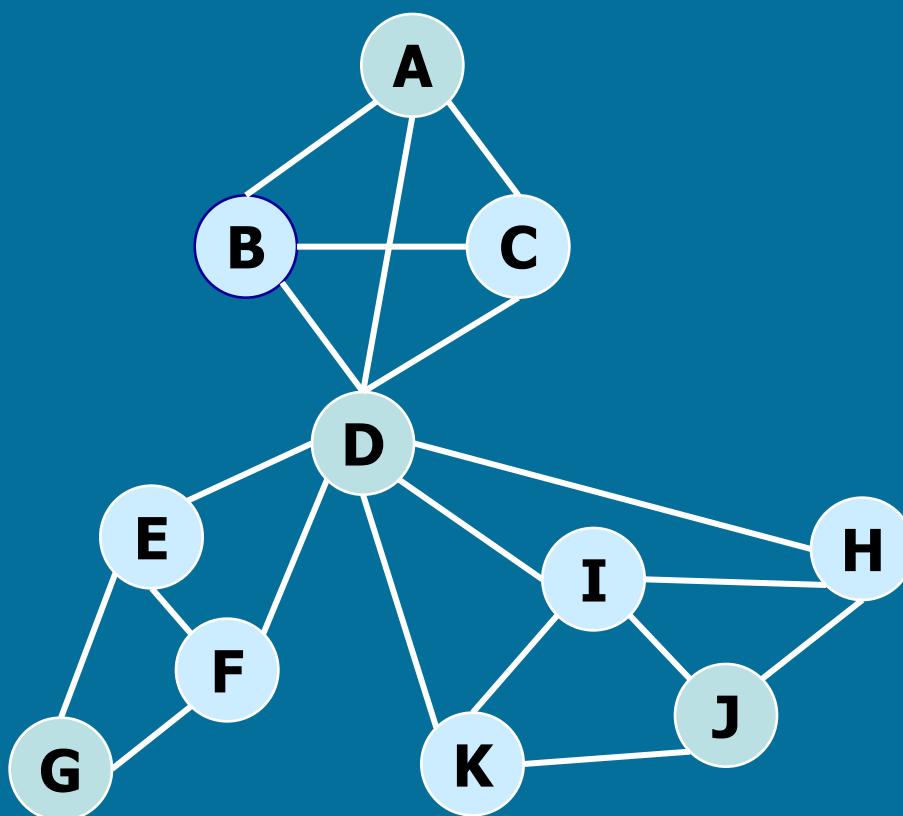
$d = 44$

$e = 1539$

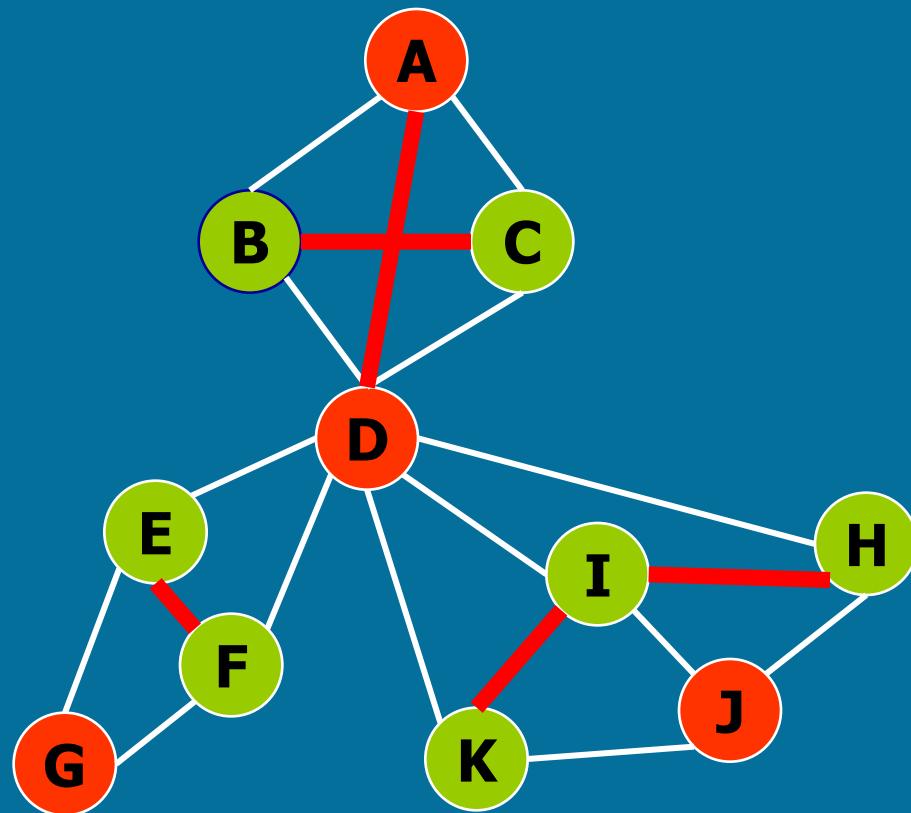
$w = 85$

$s = 4$



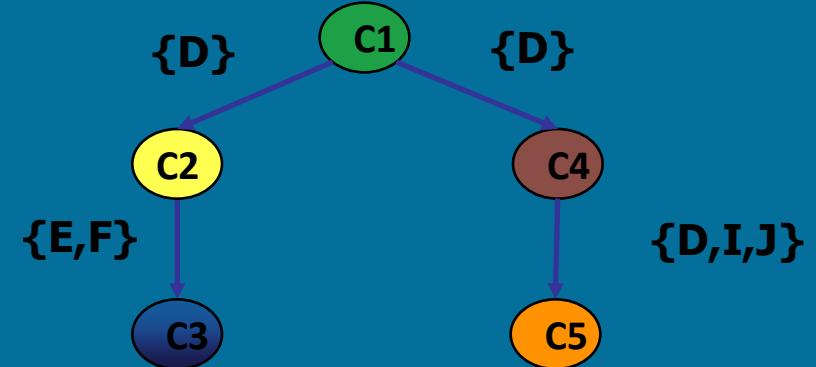
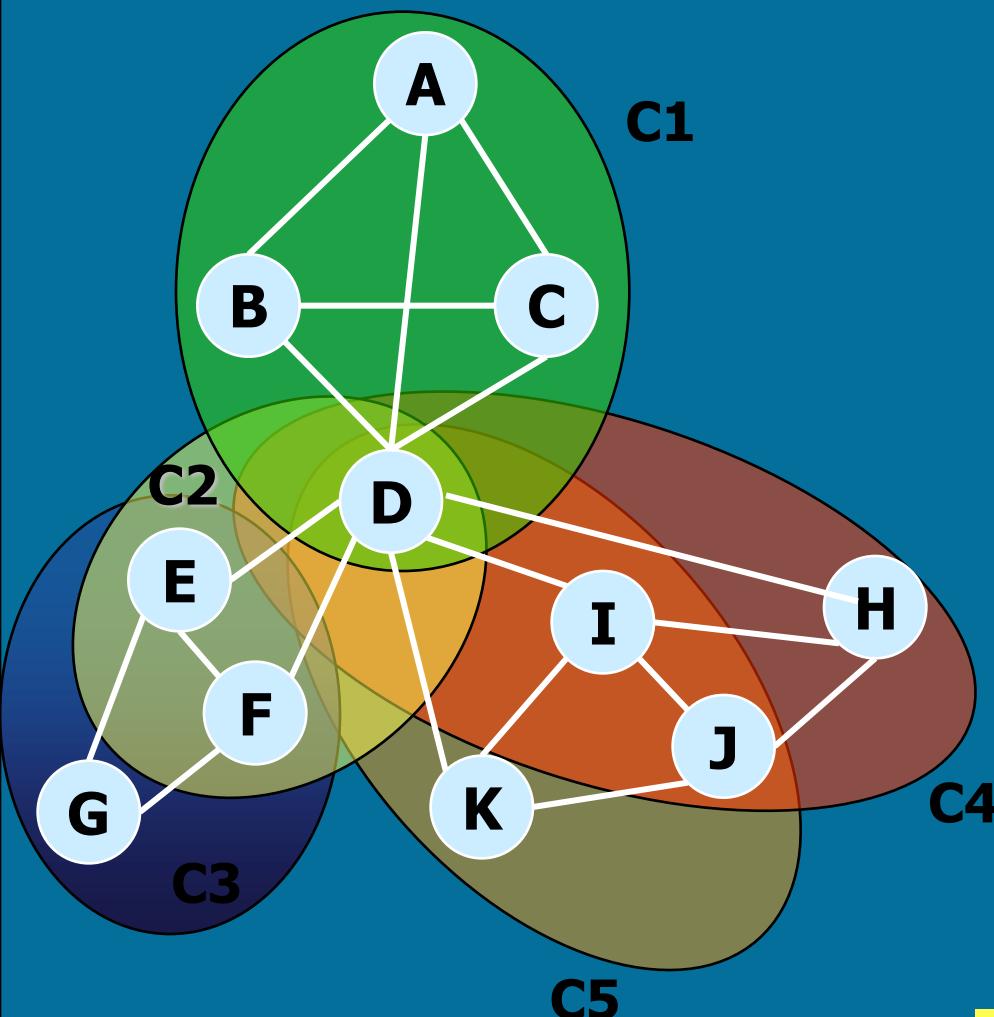


X_i	X_j	c_{ij}
1	1	1
1	0	0
0	1	0
1	1	1



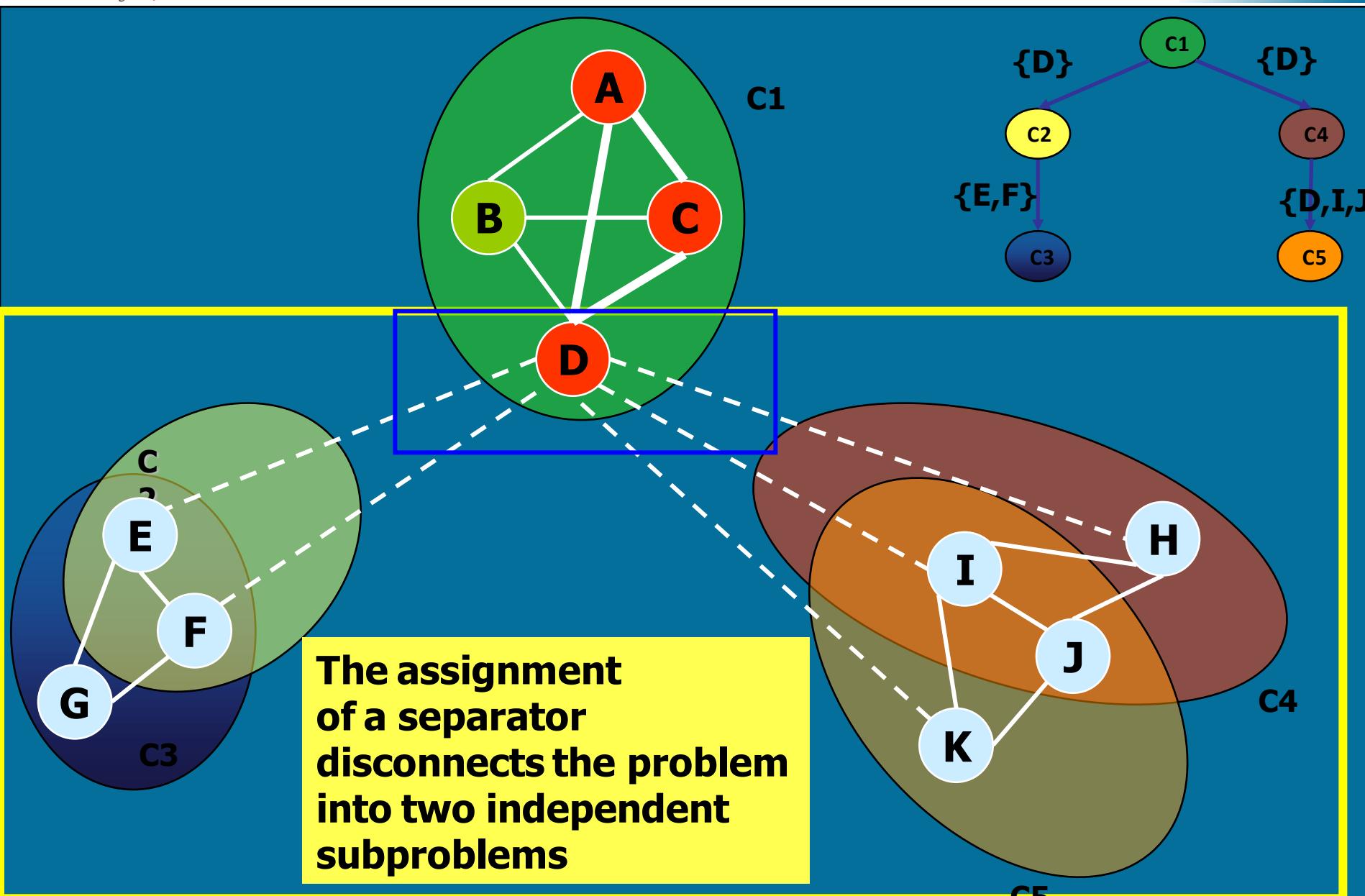
Optimal solution
with a cost of 5

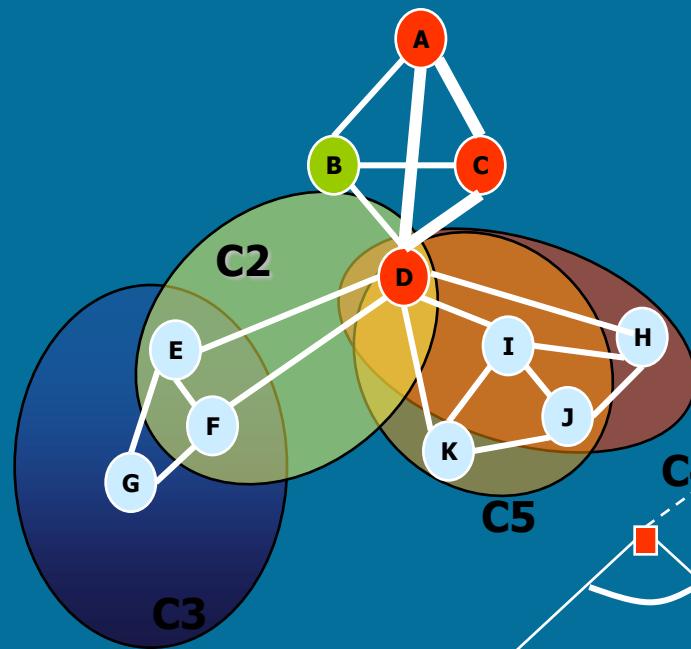
Tree Decomposition



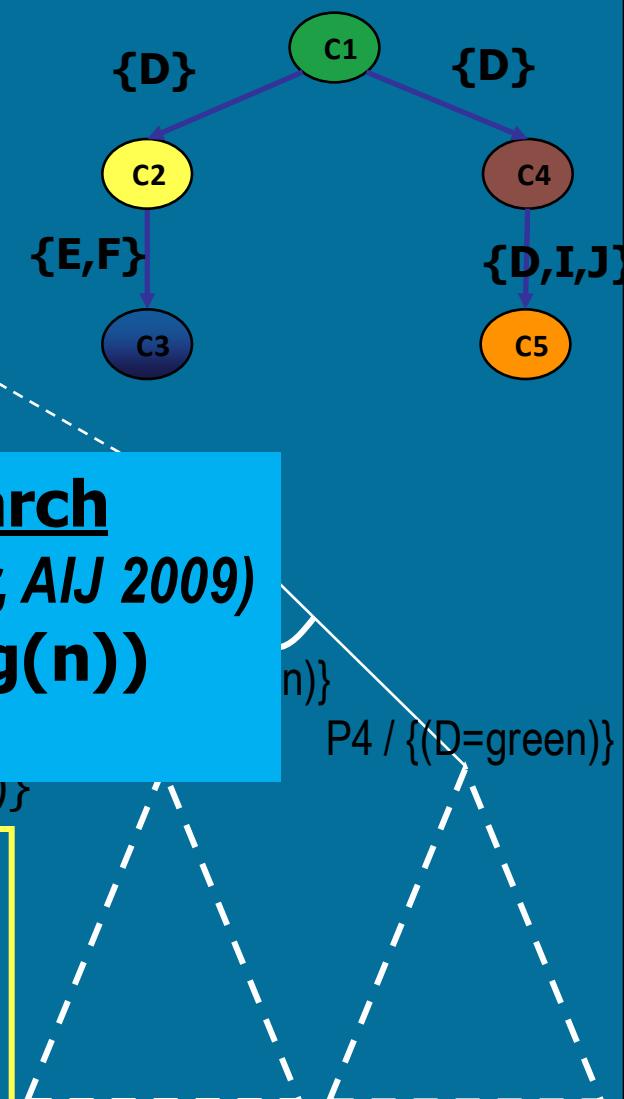
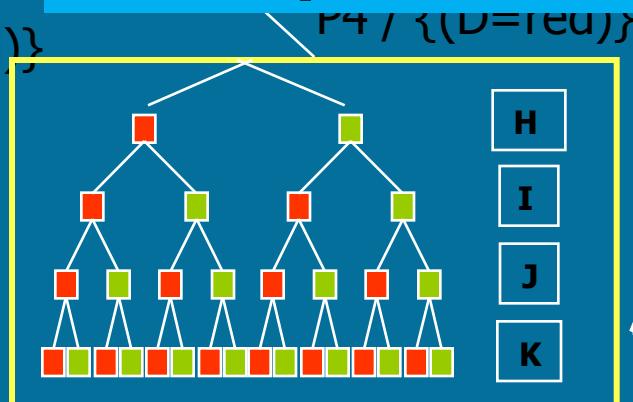
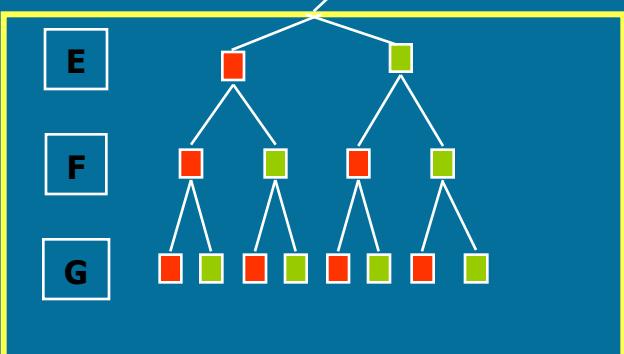
The set of clusters covers the set of variables and the set of cost functions

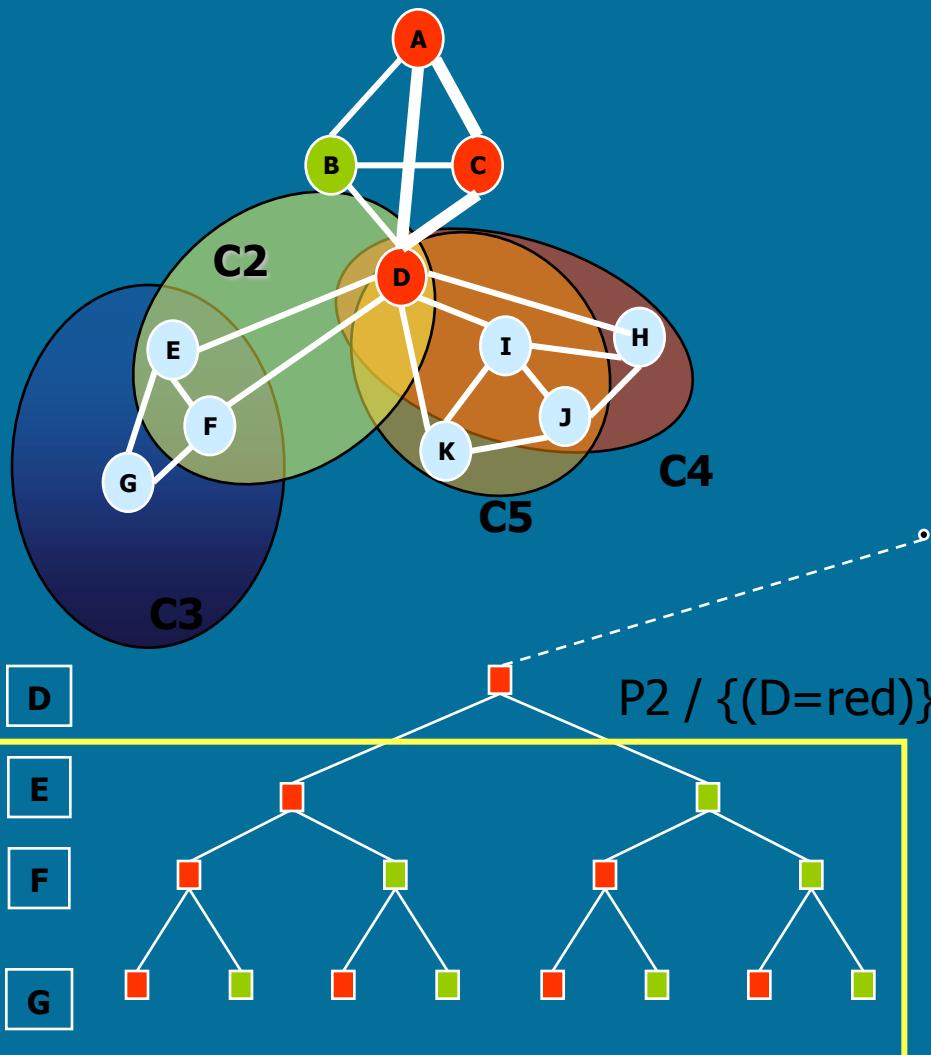
Separator = intersection between two connected clusters

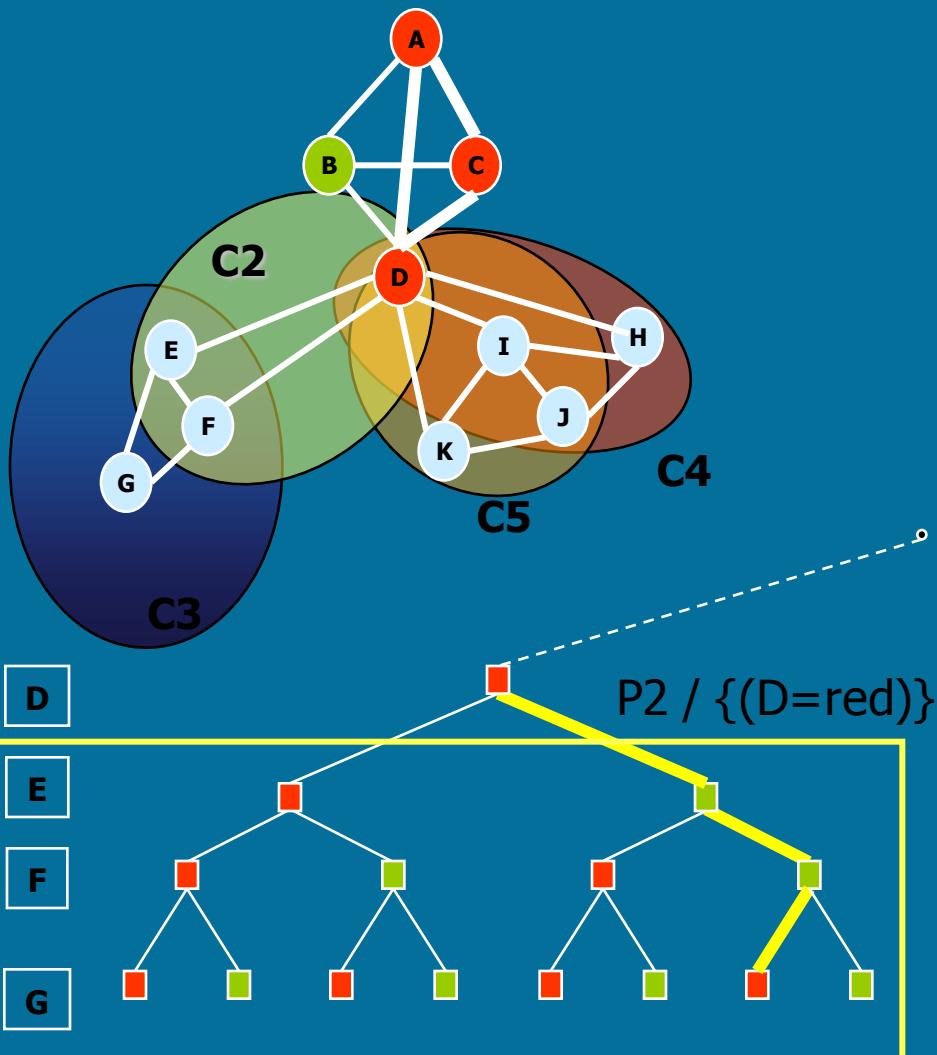




AND/OR tree search
(Marinescu & Dechter, AIJ 2009)
time $O(\exp(w \log(n)))$
linear space





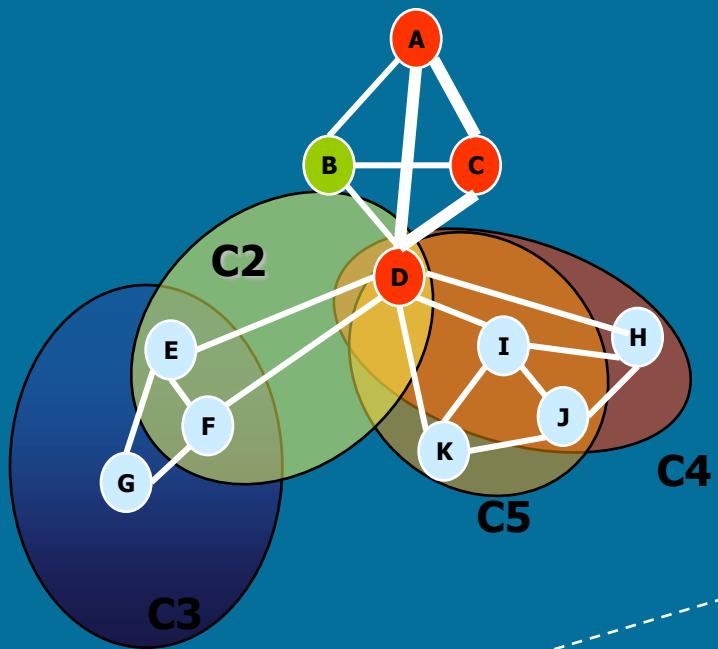


Record the optimum
of $P2 / \{(D=\text{red})\}$

AND/OR graph search
(Marinescu & Dechter, AIJ 2009)
time $O(\exp(w))$
space $O(\exp(w))$

bound $k = 5$.

**It may be useless to
compute the optimum of
 $P2 / \{(D=\text{red})\}$,
only a lower bound is
needed!**

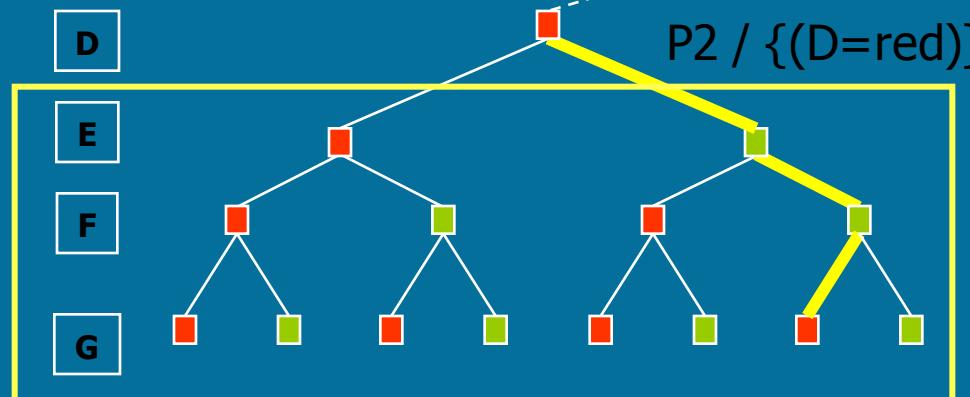


BTD

(Jégou & Terrioux, ECAI 2004)
 (de Givry et al., AAAI 2006)

time $O(k * \exp(w))$
space $O(\exp(w))$

per



It may be useless to compute the optimum of P_2 ,
 only a lower bound is needed!

Add a local upper bound:
 $UB_{P2 / \{(D=red)\}} = k - 3 - LB_{P4 / \{(D=red)\}}$

$$UB_{P2 / \{(D=red)\}} = k - 3 - \max (f_\emptyset^{C4} + f_\emptyset^{C5}, LB_{P4 / \{(D=red)\}})$$

Maintaining local consistency Recorded during search

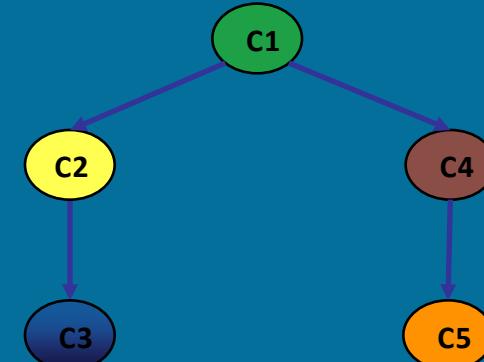
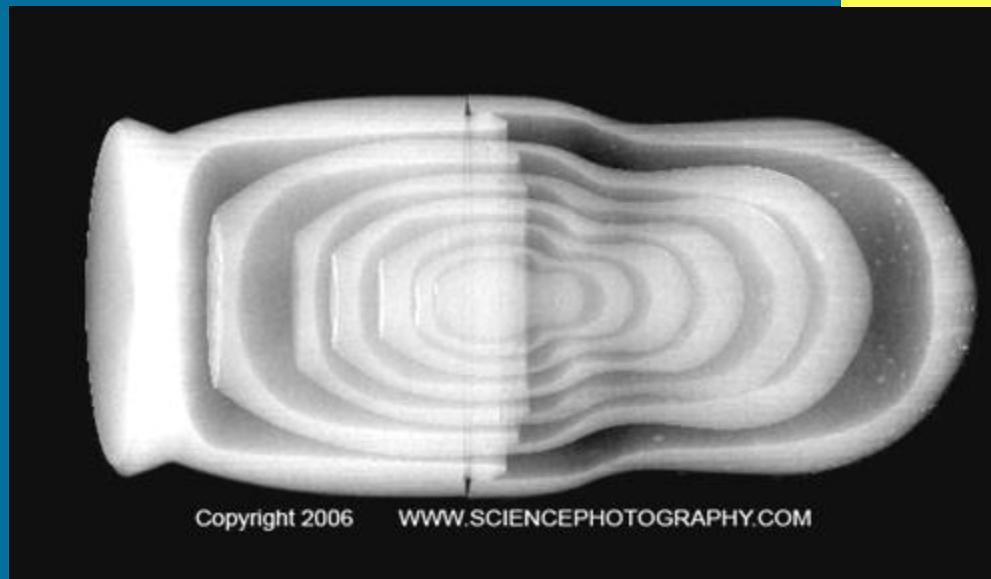
(Sanchez et al, IJCAI 2009)

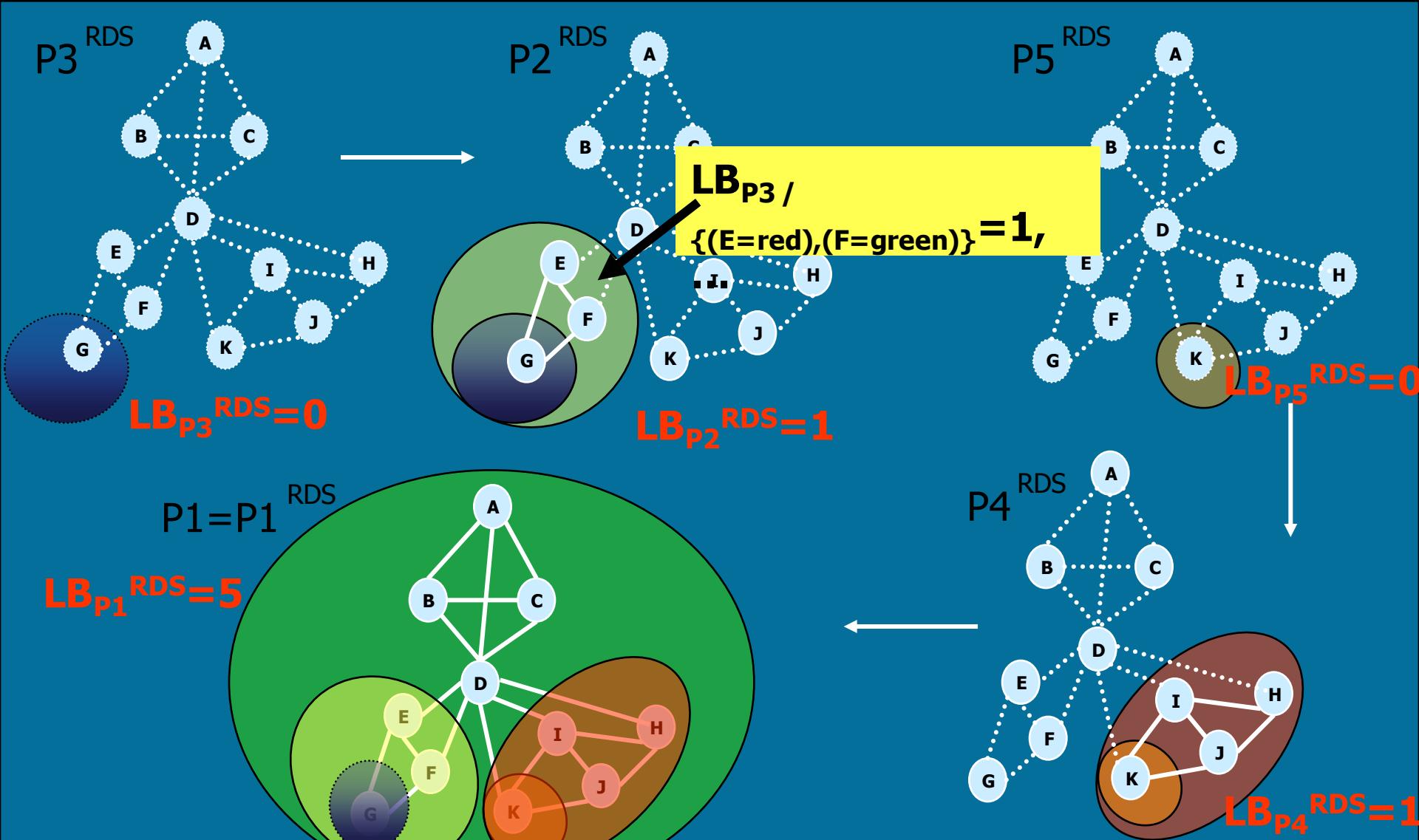
$$\text{UB}_{P2 / \{(D=\text{red})\}} = k - 3 - \max (f_{\emptyset}^{C4} + f_{\emptyset}^{C5}, \underbrace{\text{LB}_{P4 / \{(D=\text{red})\}}, \text{LB}_{P4^{\text{RDS}}} }_{\substack{\text{Found during search} \\ \text{Obtained in preprocessing}}})$$

RDS-BTD lower bound is a maximum between

- Local Consistency
- Recorded lower bounds
- RDS lower bounds

The optimum of P_i^{RDS}
is a lower bound of P_i / A
for any separator assignment A



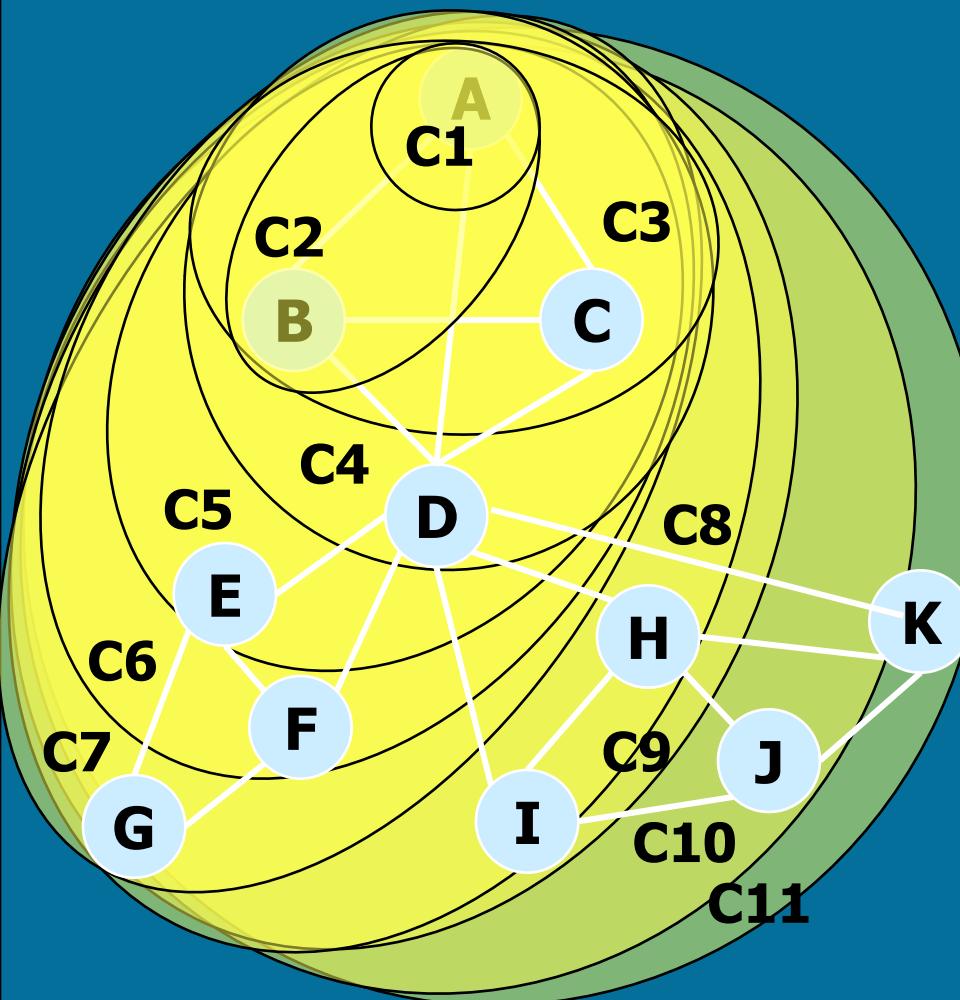


Same complexities as BTD: Time: $O(\exp(w+1))$ Space: $O(\exp(s))$

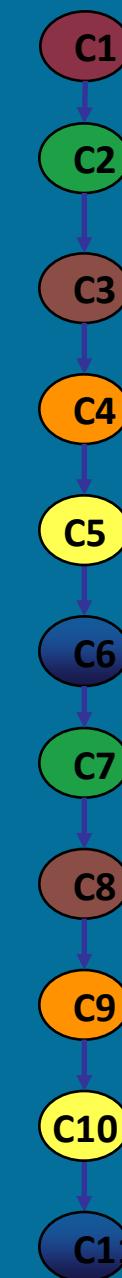
$w = \text{tree-width} = \text{largest cluster size} - 1,$

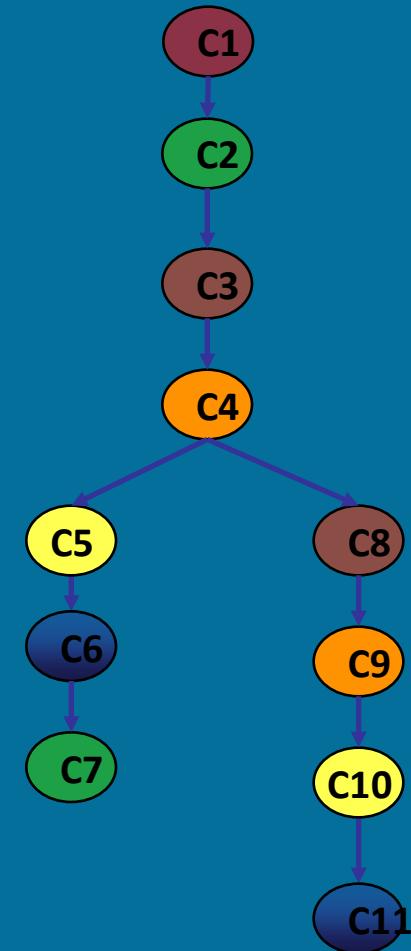
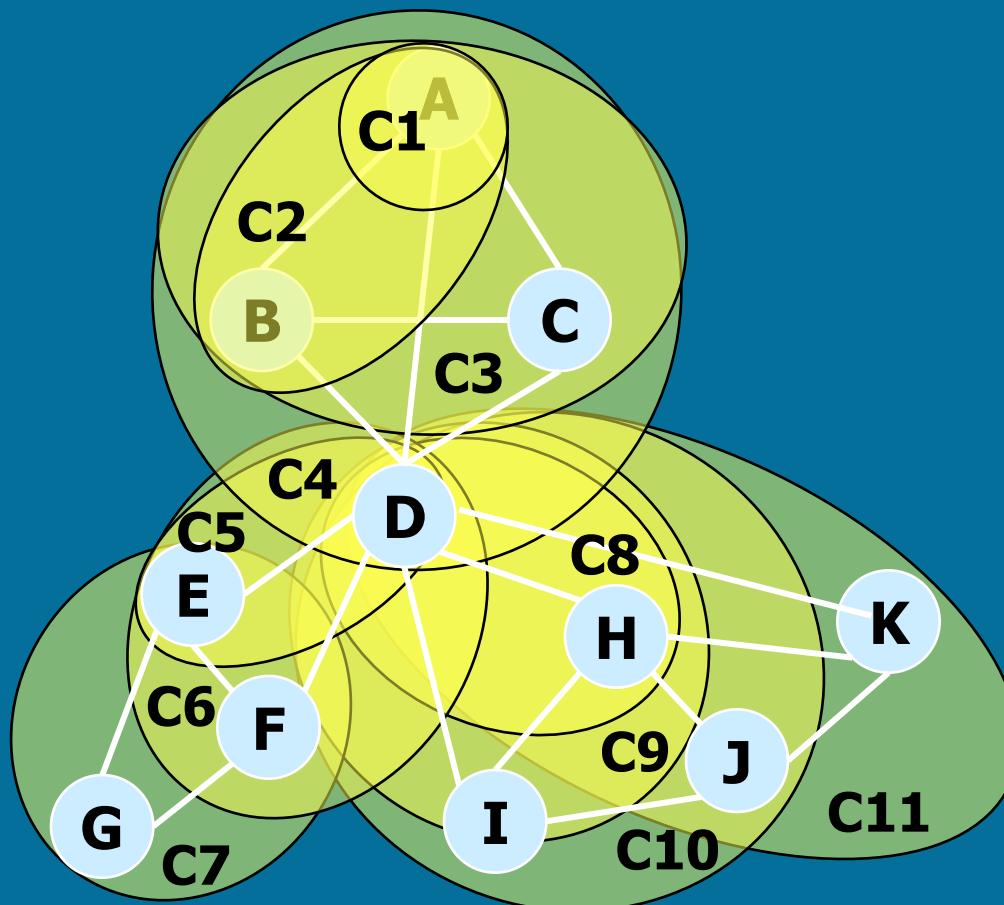
$s = \text{largest separator size}$

Related Work



RDS (Verfaillie *et al*, AAAI 1996)



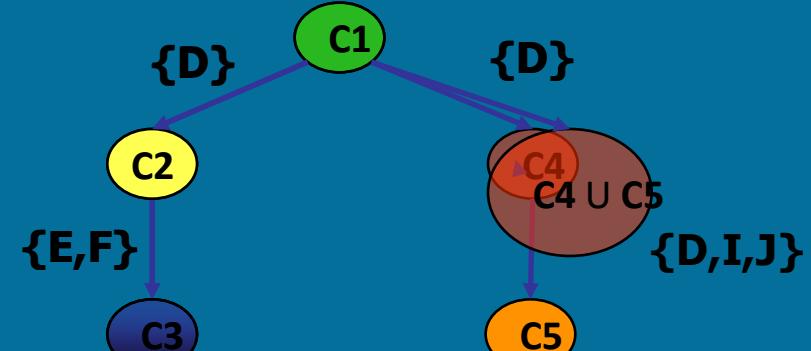
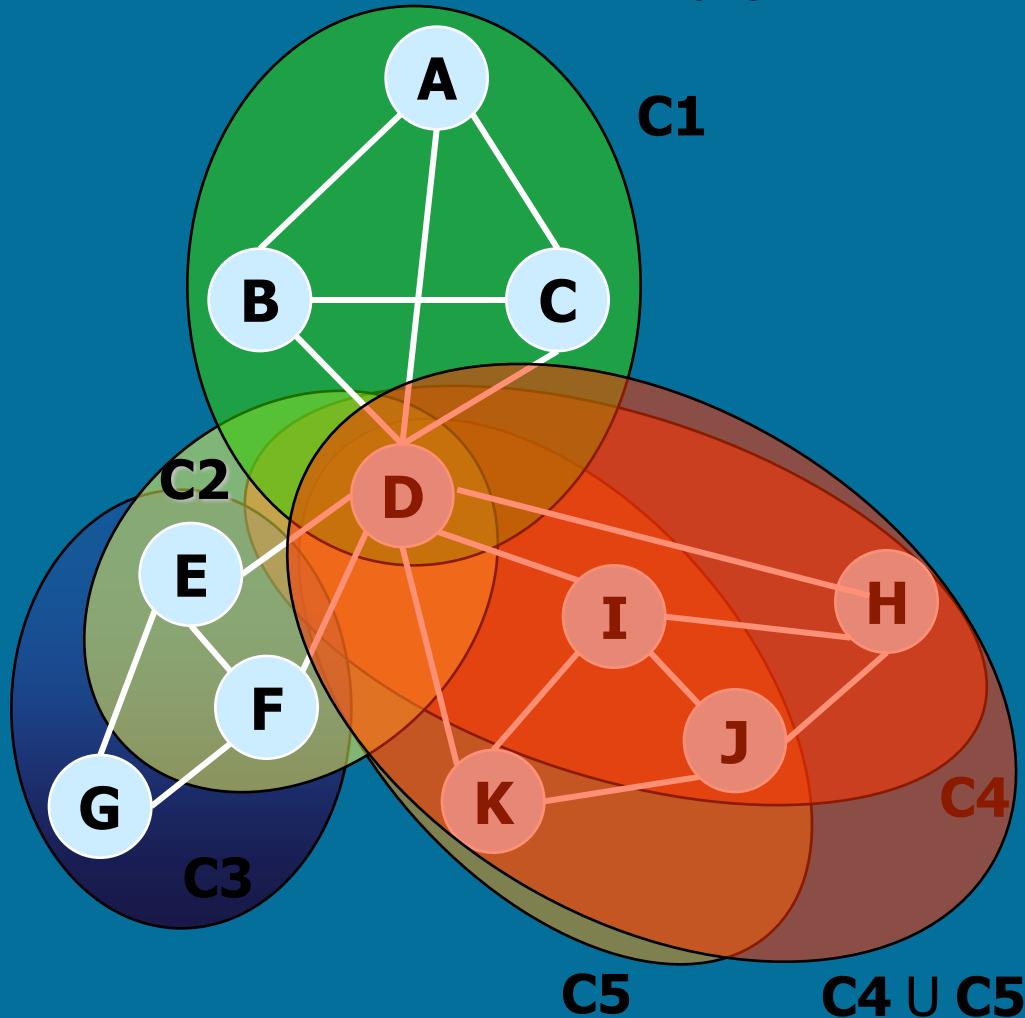


Pseudo-Tree RDS (Larrosa *et al*, ECAI 2002)

AND/OR graph search (Marinescu & Dechter, AAAI 2006)

Exploiting small separators only

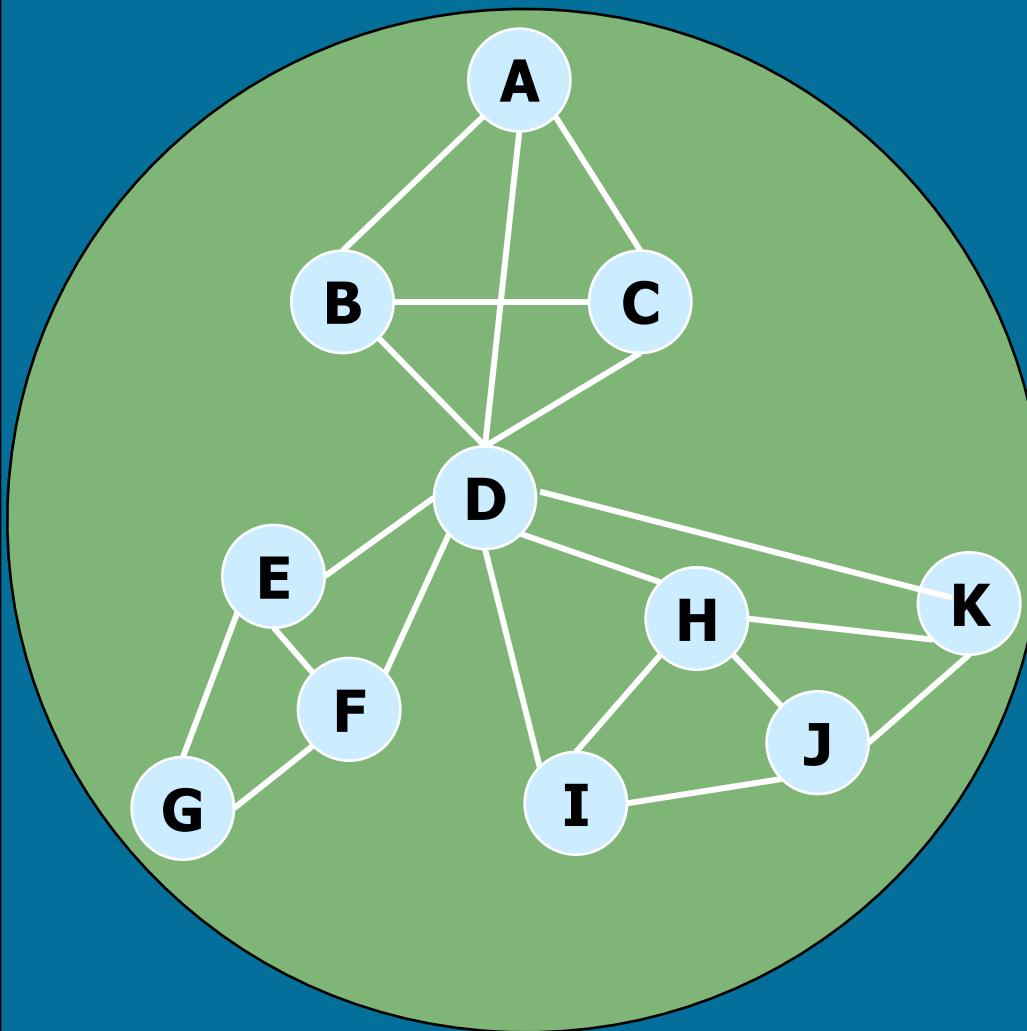
(Jégou et al., CP 2007)



$p = 2$

Merge clusters i and $\text{father}(i)$
if $|\text{sep}_{i,\text{father}(i)}| > p$

→ Gives more freedom for the dynamic variable ordering heuristic



c1

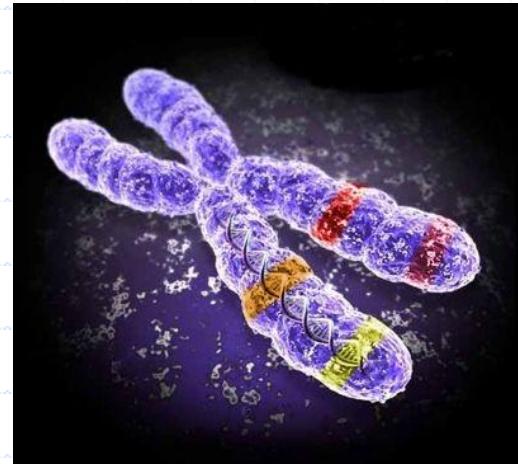
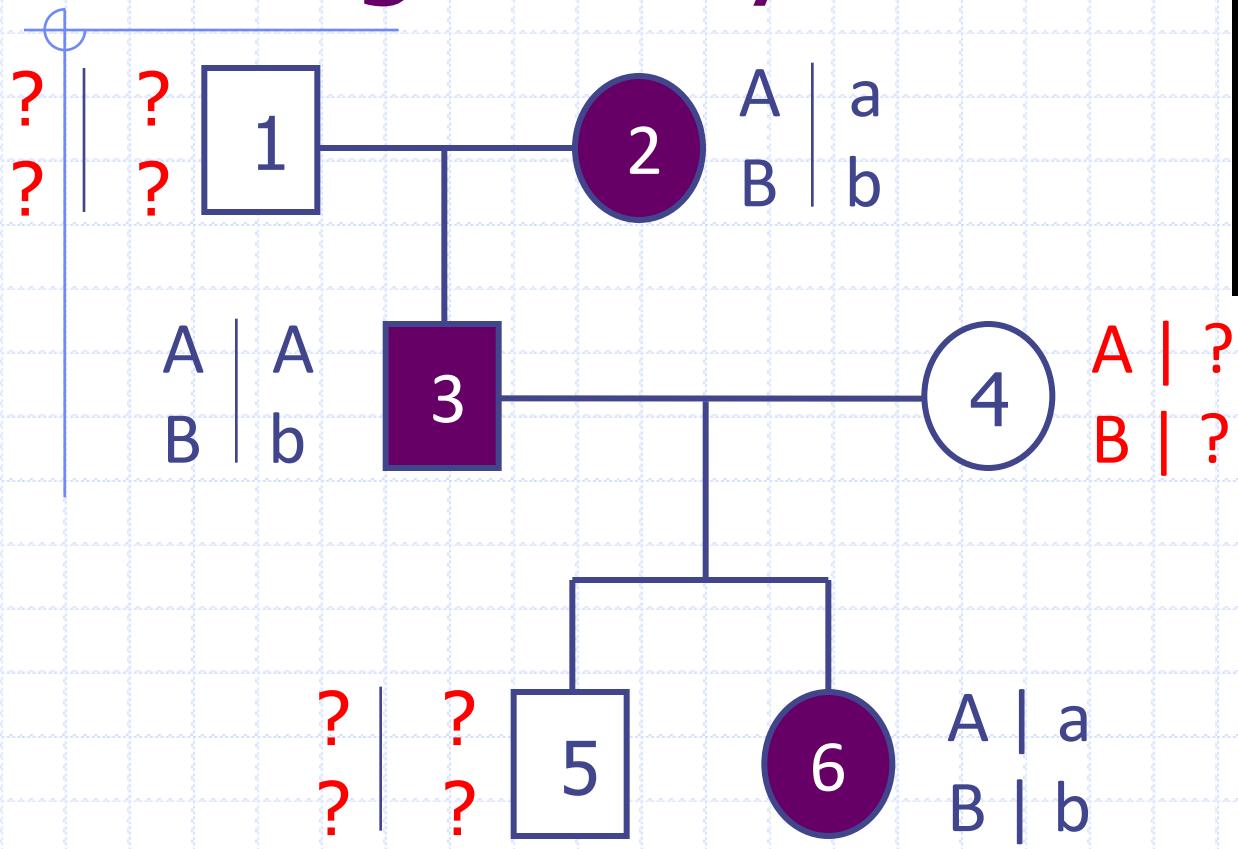
p = 0

DFBB

Some practical observations

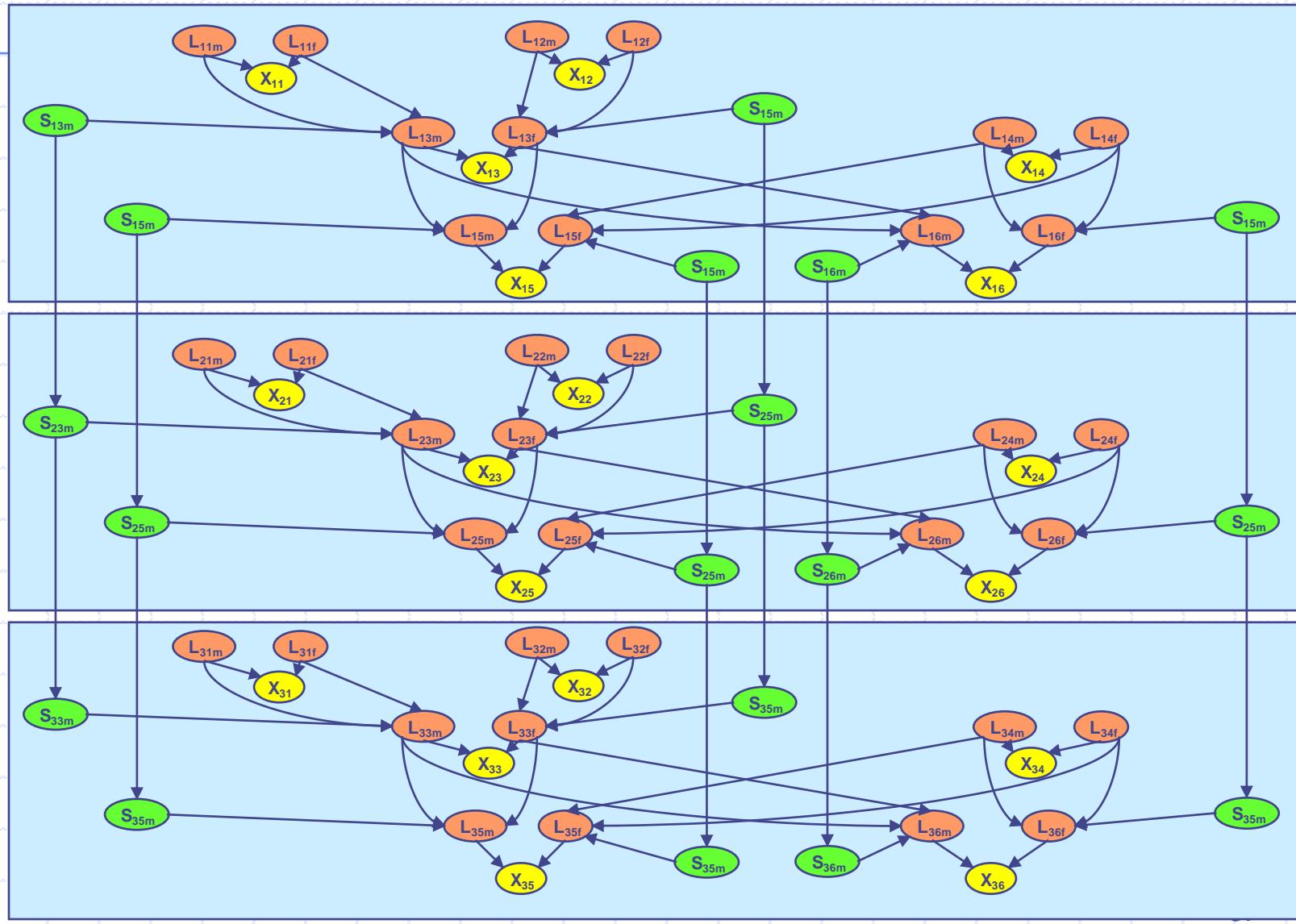
- ◆ BTD may use much less memory than Variable Elimination thanks to pruning
- ◆ Impact of root cluster
 - ⇒ Choose the largest / most costly cluster as root
- ◆ Exploit small separators only
(Jégou, Ndiaye & Terrioux, CP 2007)
 - ⇒ Give more freedom for the dynamic variable ordering heuristic
 - ⇒ Tuning based on treewidth versus separator size
- ◆ BB-VE(2) often faster than BTD

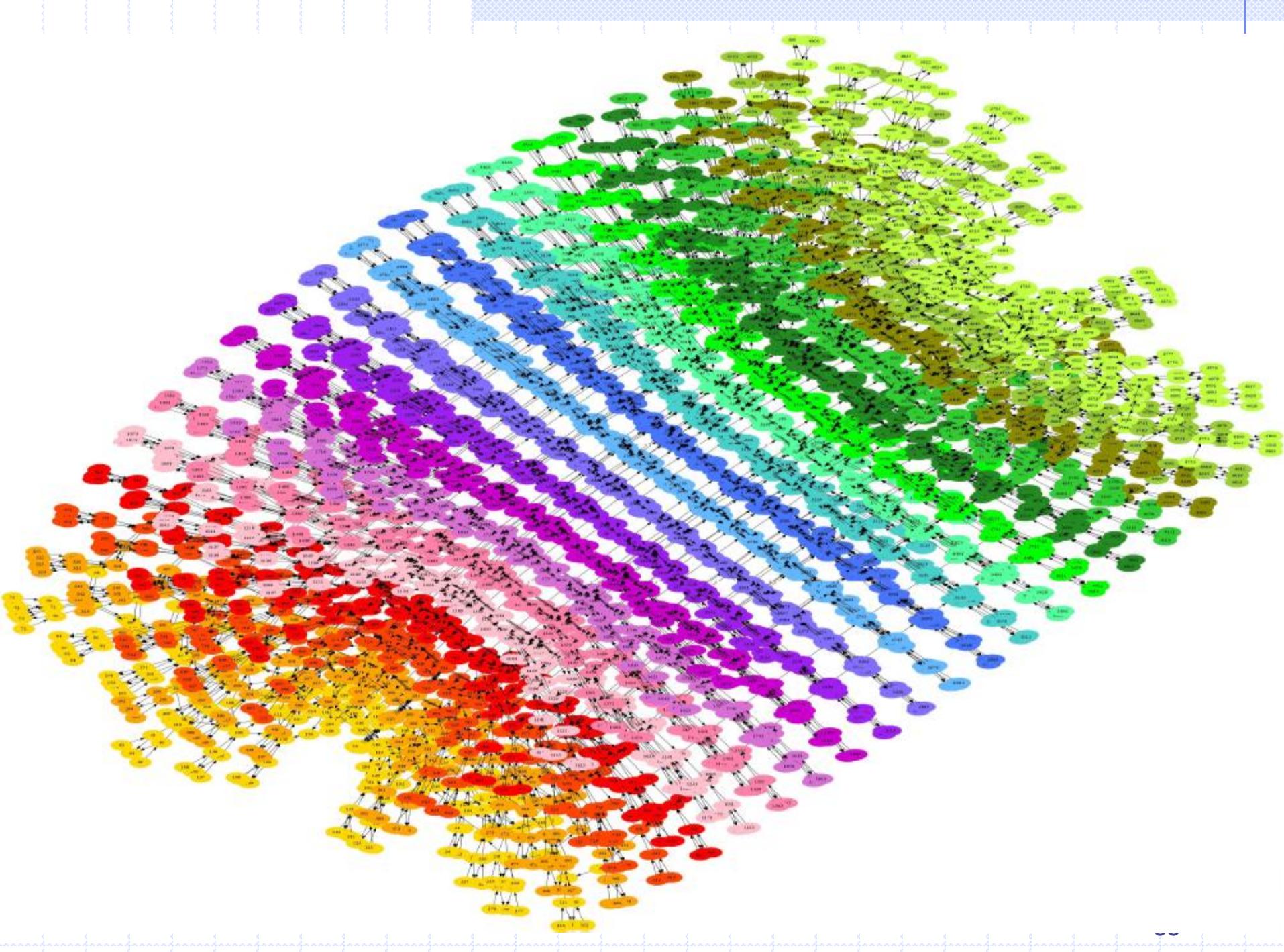
Linkage Analysis



- 6 individuals
- Haplotype: {2, 3}
- Genotype: {6}
- Unknown

Pedigree: 6 people, 3 markers





Genetic Linkage Analysis

(Fishelson et al, J. Hum. Heredity 05)

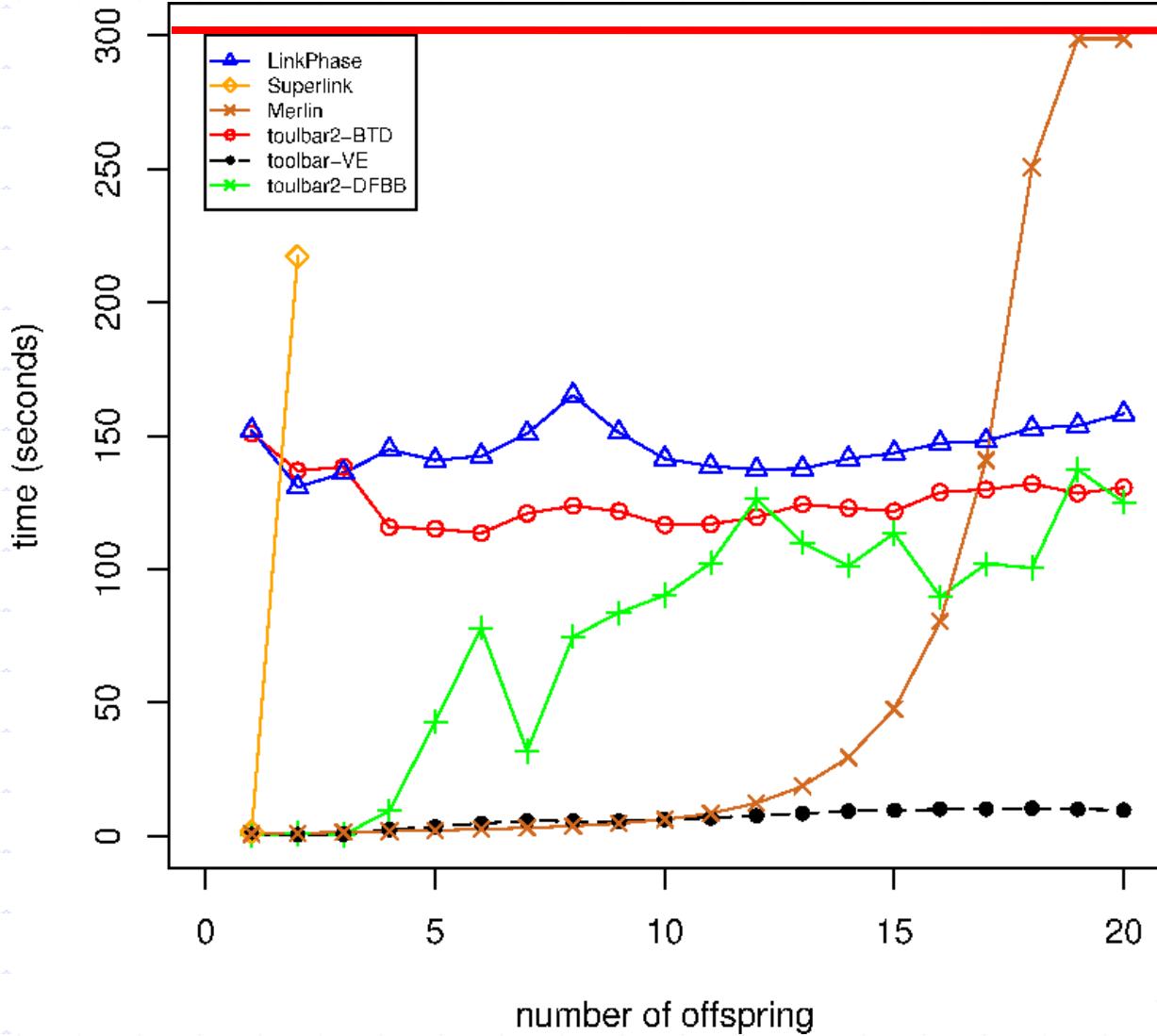
pedigree (n, d) (w*, h)	Superlink v. 1.6	SamIam v. 2.3.2	MBE(i) BB+SMB(i) AOBB+SMB(i)		MBE(i) BB+SMB(i) AOBB+SMB(i)		MBE(i) BB+SMB(i) AOBB+SMB(i)	
			i=12	time	nodes	i=16	time	nodes
ped18 (1184, 5) (21, 119)	139.06	157.05	0.51	-	-	4.59	-	-
			-	-	-	270.96	2,555,078	20.27
ped25 (994, 5) (29, 53)	-	out	0.34	-	-	3.20	-	-
			-	-	-	-	-	-
ped30 (1016, 5) (25, 51)	13095.83	out	0.31	-	-	2.66	-	-
			5563.22	63,068,960	1811.34	20,275,620	82.25	588,558
ped33 (581, 5) (26, 48)	-	out	0.41	-	-	5.28	-	-
			2335.28	32,444,818	62.91	807,071	76.47	320,279
ped39 (1272, 5) (23, 94)	322.14	out	0.52	-	-	8.41	-	-
			-	-	4041.56	52,804,044	141.23	407,280

Min-fill pseudo tree. Time limit 3 hours.

Haplotype reconstruction in half-sib pedigrees

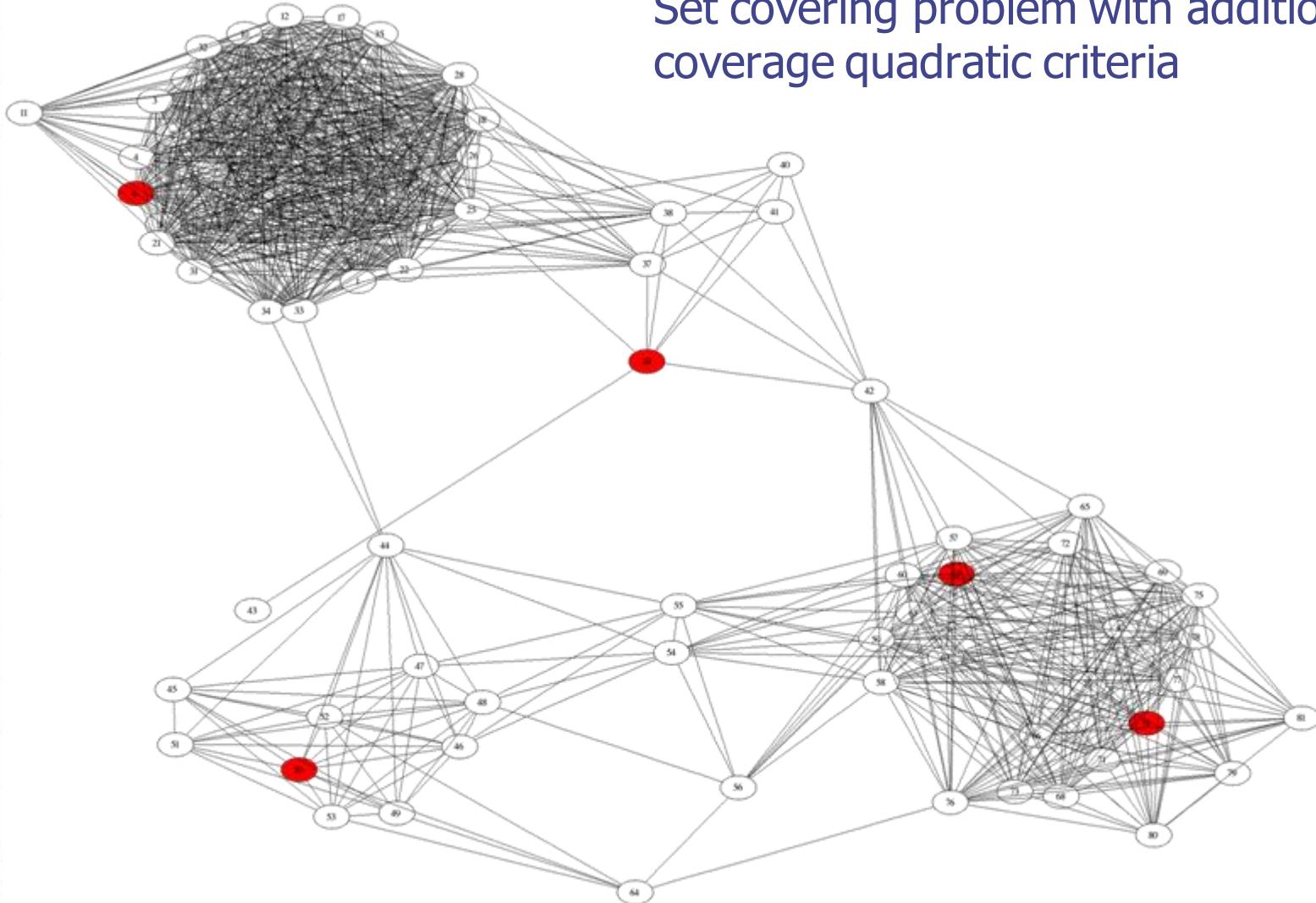
(Favier et al, WCB'10)

Read dataset
HAPMAP
Chr.X with
36,000 markers
treewidth < 15



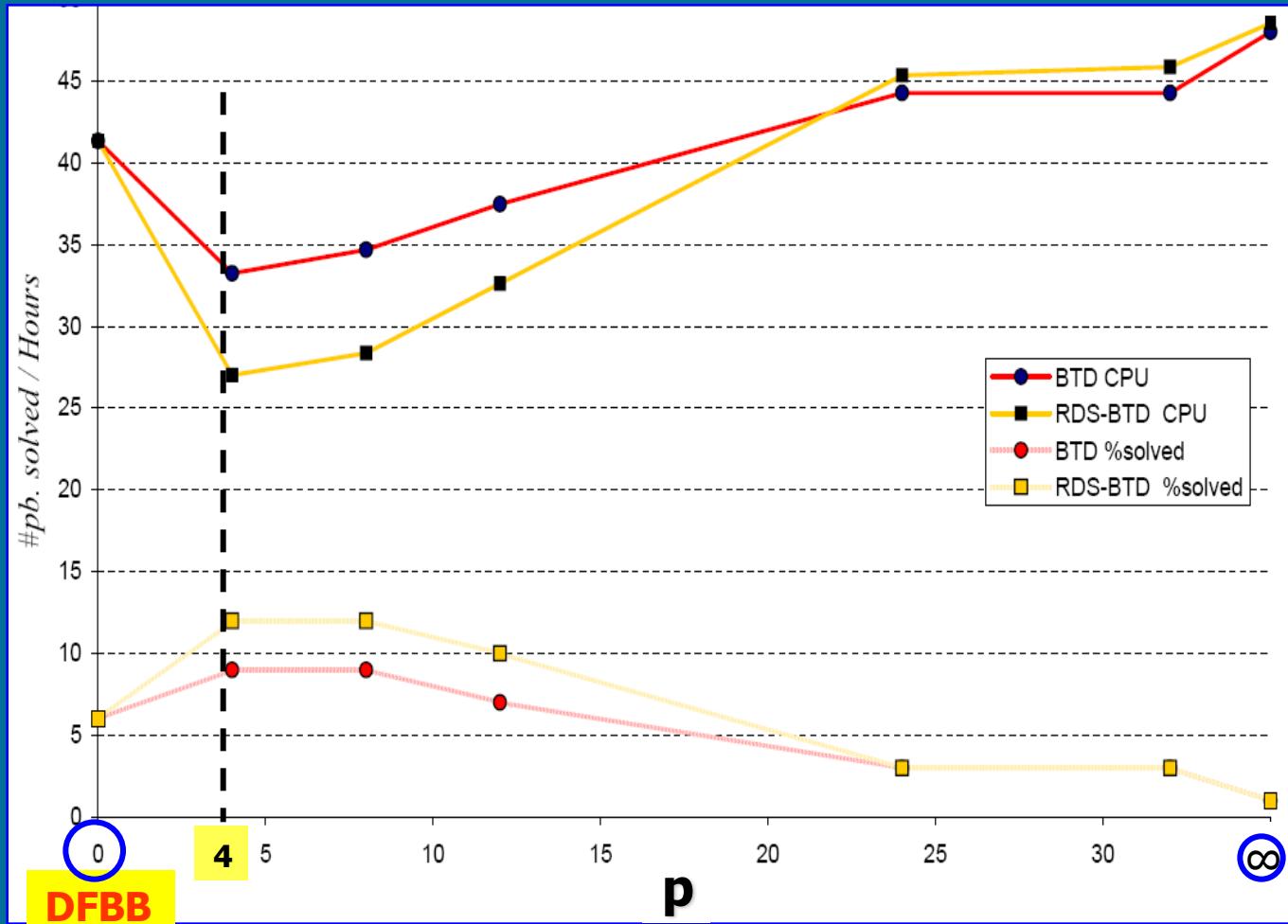
Tag SNP Selection

Set covering problem with additional coverage quadratic criteria



Tag SNP Selection

25 largest instances from HapMap human chr01 ($r^2 \geq 0.5$)



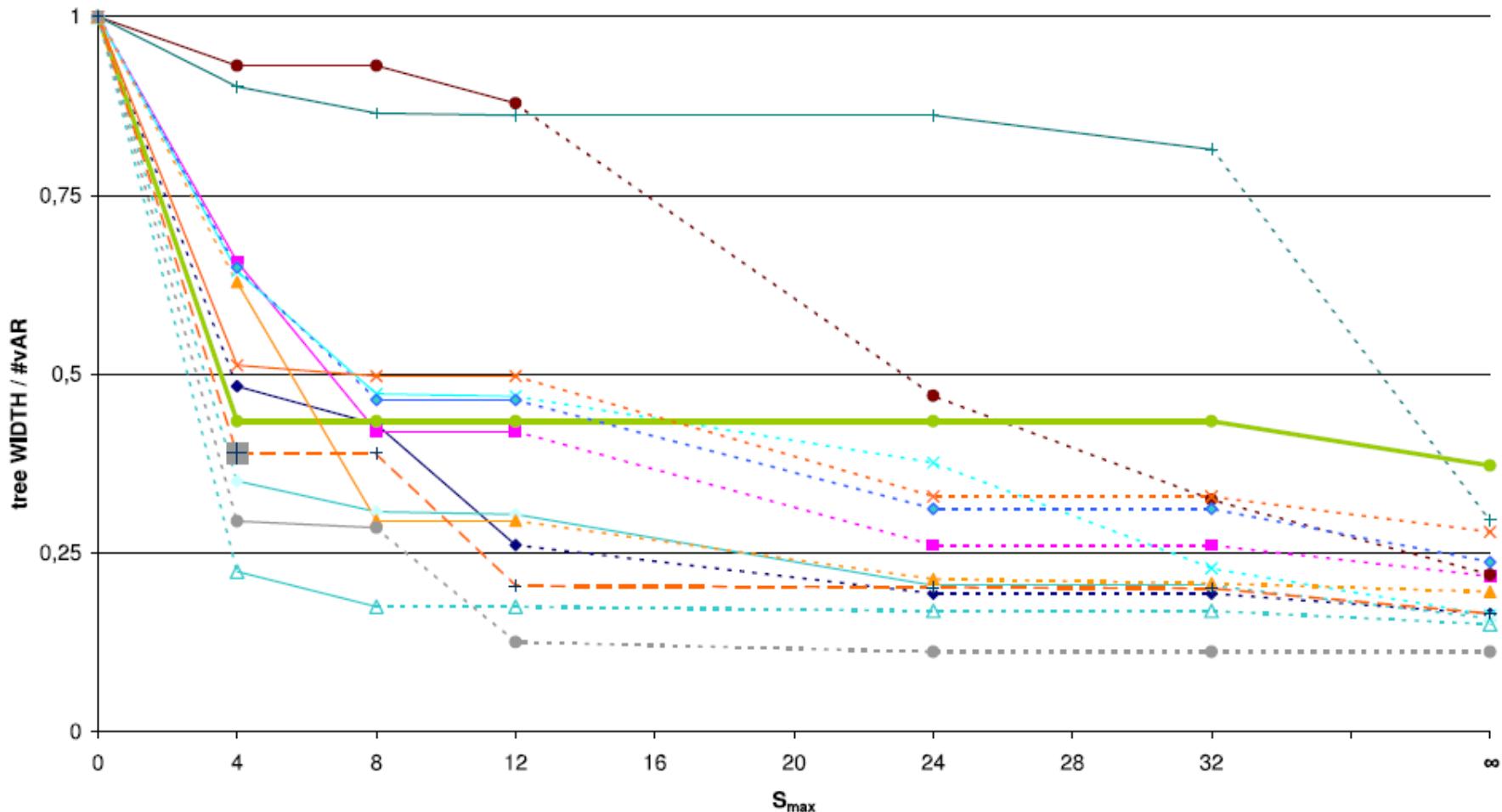
$171 \leq n/2 \leq 777$
 $30 \leq d \leq 266$
 $6\% \leq m \leq 37\%$
 $14\% \leq w \leq 23\%$

Solved by
toulbar2 v.0.8
with initial
upper bound
found by FESTA
and with MCS
tree decomposition

Time limit: 2 hours

BTD
RDS-BTD

Tuning the separator size limit for the Tag SNP problem (Allouche et al., WCB'09)



Evolution of treewidth normalized by variable number
for different maximum separator size thresholds

Bibliography

- ◆ For hybrids of search and inference, see the chapter 10 in *Constraint Processing*, Dechter, Morgan Kaufmann, 2003.
- ◆ For exploiting tree decomposition, see
 - “*Exploiting Tree Decomposition and Soft Local Consistency in Weighted CSP*”, de Givry, Schiex & Verfaillie , AAAI 2006.
 - “*Memory intensive AND/OR search for combinatorial optimization in graphical models (Part I&II)*”, Marinescu & Dechter, AIJ 2009.

Applications / benchmarks

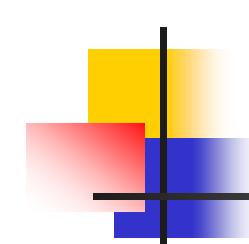
toulbar2, aolib

◆ Resource Allocation

- Frequency assignment (Allouche et al, CP 2010) CTE, BTD, VAC
 $n \leq 458$, $d=44$, $e(2) \leq 5,000$
- Satellite management (Verfaillie et al, AAAI 1996) RDS, RDS-BTD
 $n \leq 364$, $d=4$, $e(2-3) \leq 10,108$
- Uncapacitated warehouse location (Zytnicki et al, IJCAI 2005) EDAC, VAC,
 $n \leq 1,100$, $d \leq 300$, $e(2) \leq 100,000$ ILPO/1

◆ Bioinformatics

- Genetic linkage analysis (Marinescu & Dechter, AAAI 2006) AND/ORsearch
 $n \leq 1,200$, $d \leq 7$, $e(2-5) \leq 2,000$
- Mendelian error detection (Sanchez et al, Constraints 2008) EDAC3, BB-VE
 $n \leq 20,000$, $d \leq 66$, $e(3) \leq 30,000$
- RNA gene finding (Zytnicki et al, Constraints 2008) BAC
 $n \approx 20$, $d > 100$ million, $e(4) \approx 10$
- Tag SNP selection (Sanchez et al, IJCAI 2009) RDS-BTD, ILPO/1
 $n \leq 1,500$, $d \leq 266$, $e(2) \leq 150,000$



CP'06 Competitors

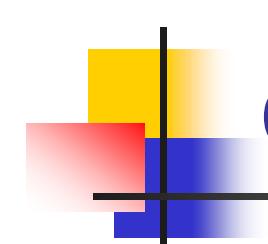
- Solvers
 - AbsconMax (ie, DFBB+MRDAC)
 - **aolibdvo** (ie, AOBB+EDAC+DVO solver)
 - **aolibpvo** (ie, AOBB+EDAC+PVO solver)
 - CSP4J-MaxCSP
 - **Toolbar** (ie, DFBB+EDAC)
 - **Toolbar_BTD** (ie, BTD+EDAC+VE)
 - **Toolbar_MaxSAT** (ie, DPLL+specific EPT rules)
 - **Toulbar2** (ie, DFBB+EDAC+VE+LDS)

CP'06 Results

Overall ranking on all selected competition benchmarks

Solver Name		Progress					
AbsconMax 109 EPFC		done 1069					
AbsconMax 109 PFC		done 1069					
aolibdvo 2007-01-17		done 821					
aolibpvo 2007-01-17		MOPT 495	SAT 26	MSAT 42	?	259	
CSP4J - MaxCSP 2006-12-19		done 1069					
toolbar 2007-01-12		MOPT 821	SAT 26	MSAT 93	?	61	
Toolbar_BTD 2007-01-12		done 821					
Toolbar_MaxSat 2007-01-19		MOPT 646	SAT 26	?	587	149	ERR 6
Toulbar2 2007-01-12		done 821					

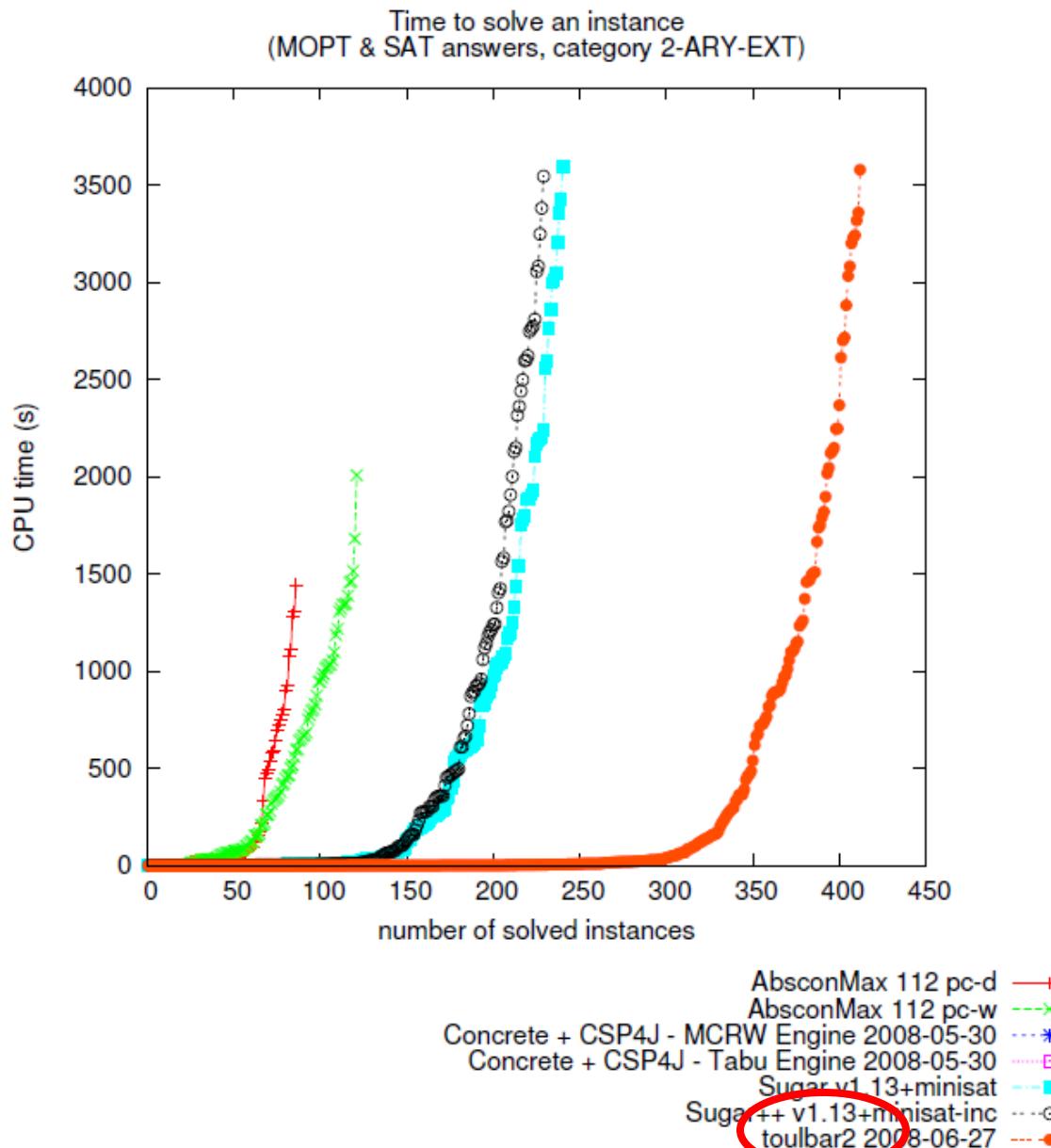
The longest dark green bar wins



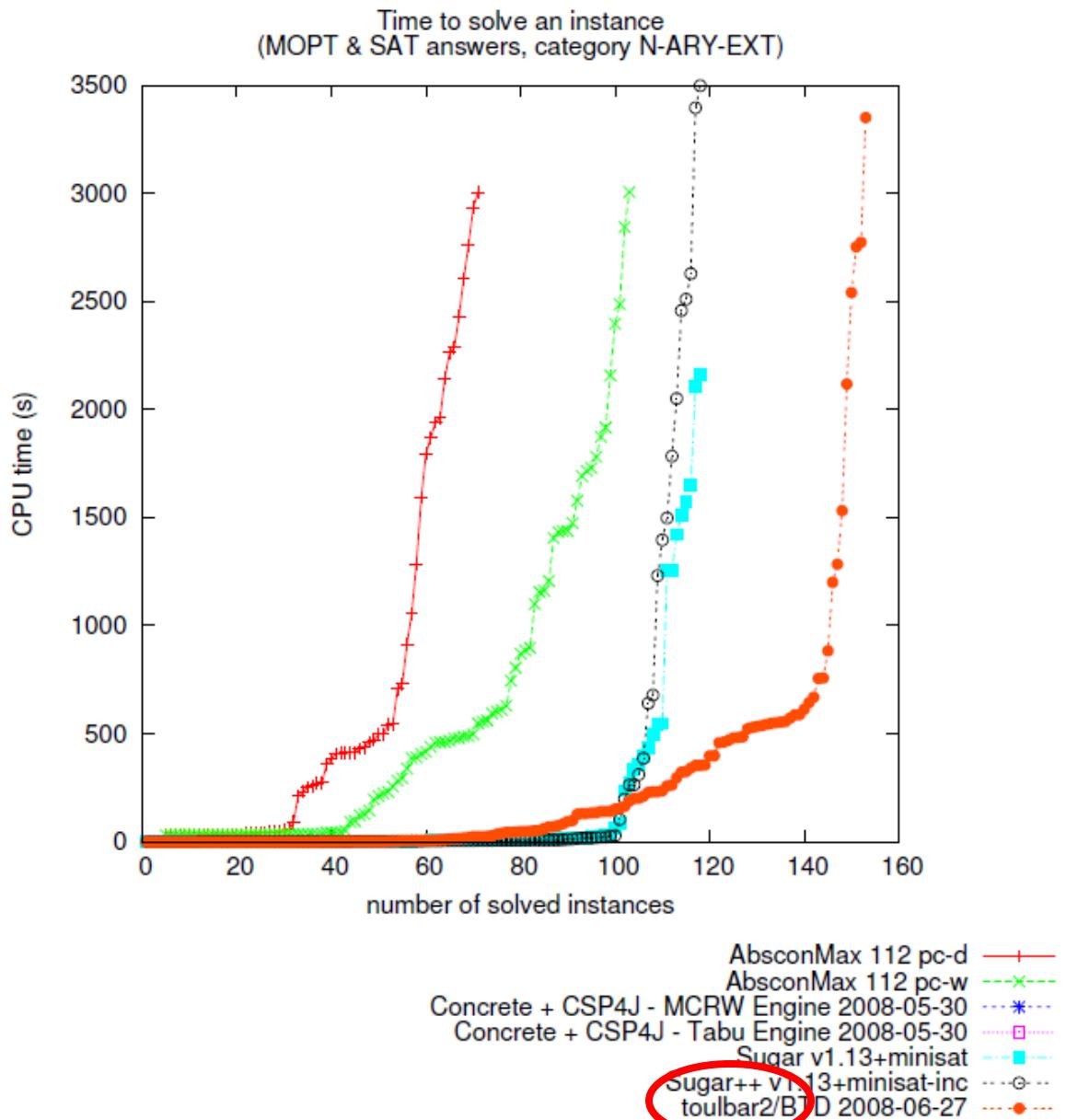
CP'08 Competitors

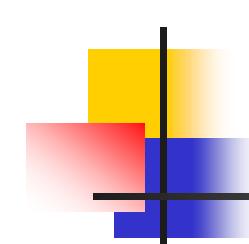
- Solvers
 - AbsconMax (ie, DFBB+MRDAC)
 - CSP4J-MaxCSP
 - Sugar (SAT-based solver)
 - **Toulbar2** (ie, BTD+EDAC+VE)

CP'08 Results



CP'08 Results

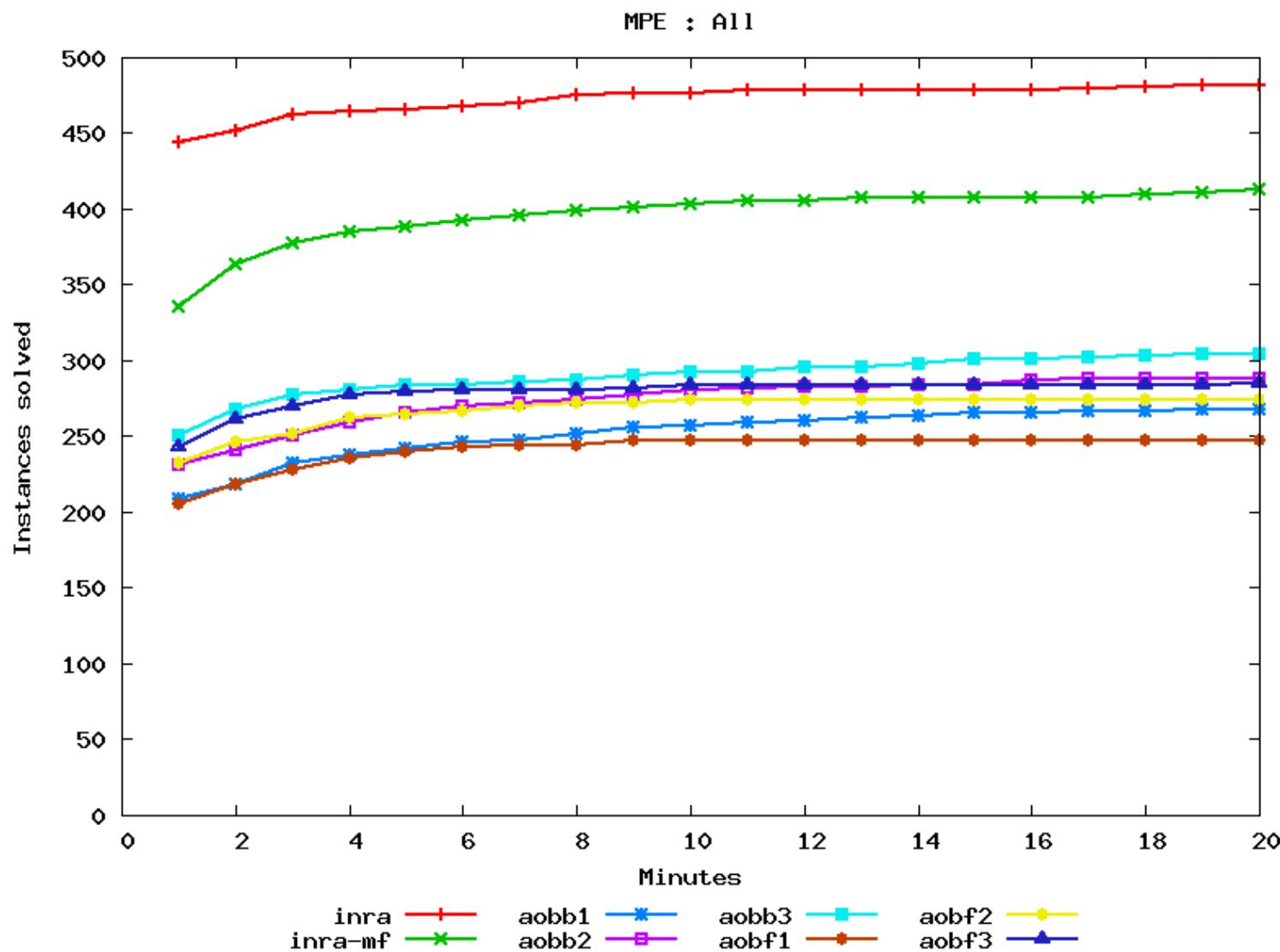




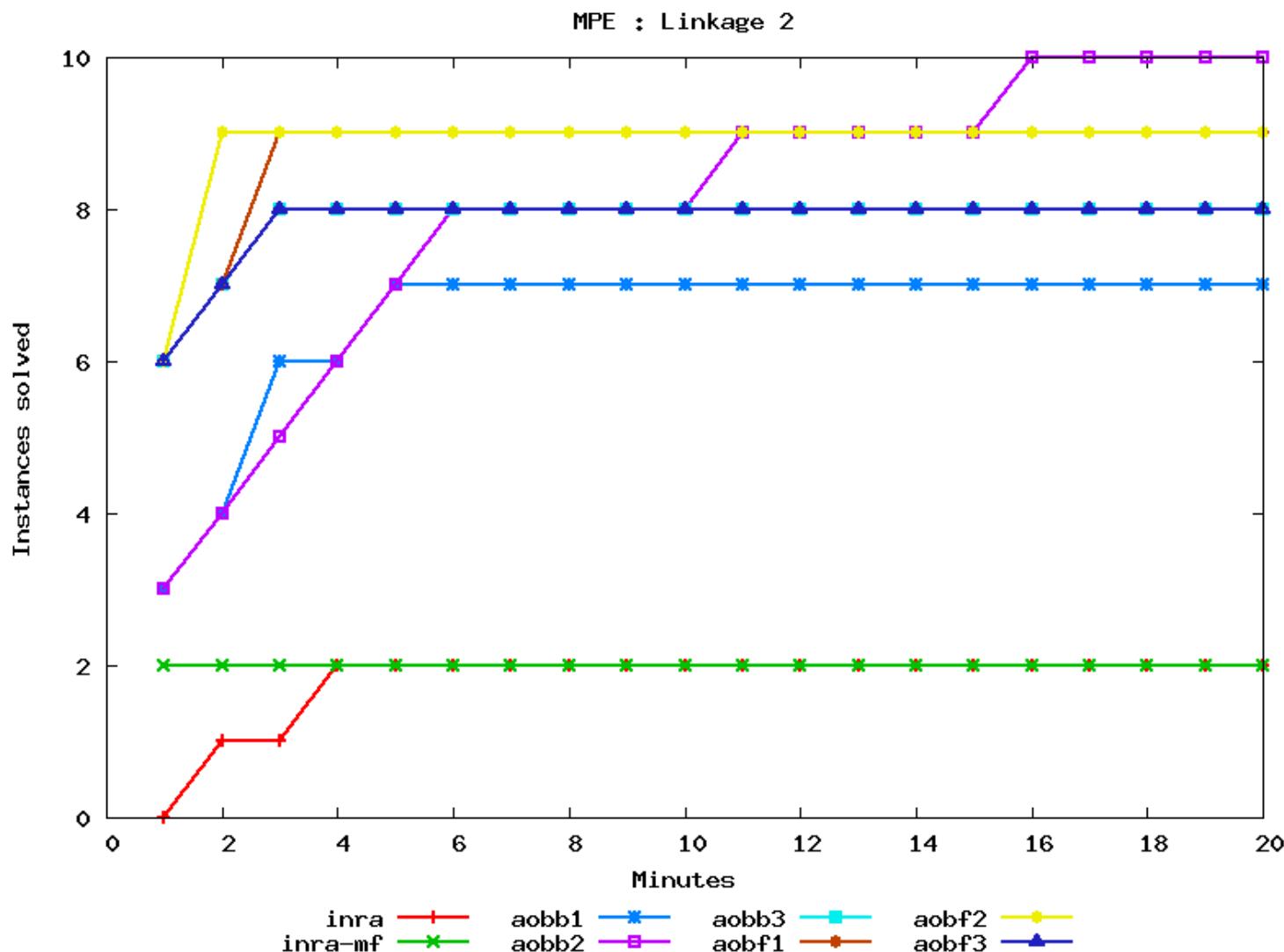
UAI'08 Competition

- **AOBB-C+SMB(i) – (i = 18, 20, 22)**
 - AND/OR Branch-and-Bound with pre-compiled mini-bucket heuristics (i-bound), full caching, static pseudo-trees, constraint propagation
- **AOBF-C+SMB(i) – (i = 18, 20, 22)**
 - AND/OR Best-First search with pre-compiled mini-bucket heuristics (i-bound), full caching, static pseudo-trees, no constraint propagation
- **Toulbar2**
 - OR Branch-and-Bound, dynamic variable/value orderings, EDAC consistency for binary and ternary cost functions, variable elimination of small degree (2) during search
- **Toulbar2/BTD**
 - DFBB exploiting a tree decomposition (AND/OR), same search inside clusters as toulbar2, full caching (no cluster merging), combines RDS and EDAC, and caching lower bounds

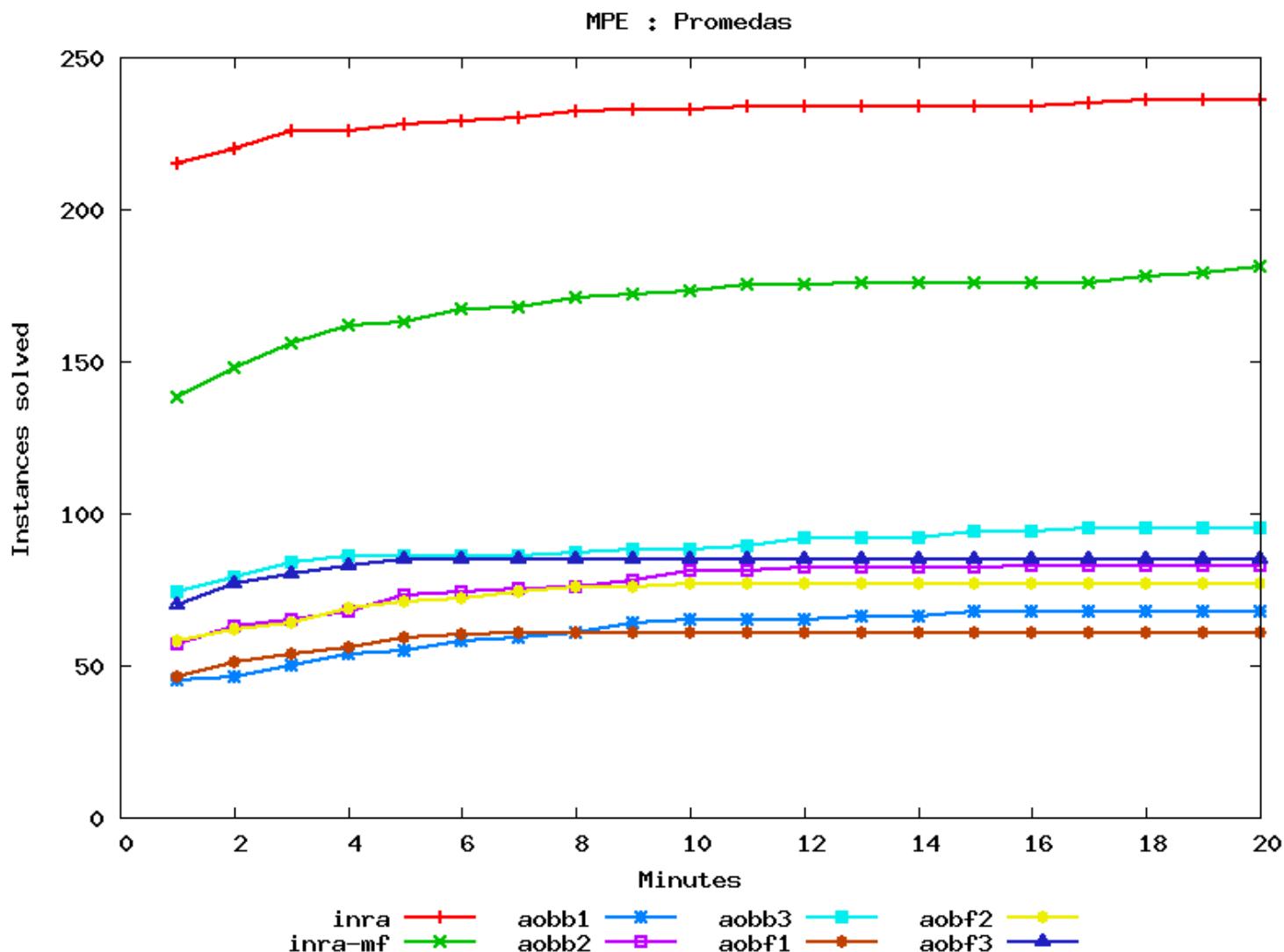
UAI'08 Competition Results



UAI'08 Competition Results (II)



UAI'08 Competition Results (III)



2010 Approximate Inference Evaluation (results given at UAI'10)

Networks – by domain (1 hour)

Network	PR	MAR	MPE
CSP	8	8	55
Grids	20	20	40
Image Alignment			10
Medical Diagnosis	26	26	
Object Detection	96	96	92
Pedigree	4	4	
Protein Folding			21
Protein-Protein Interaction			8
Segmentation	50	50	50

Summary of the results

Seconds	PR	MAR	MPE
20	Arthur Choi (UCLA)	Arthur Choi (UCLA)	Joris Mooij (Max Planck)
1200	Vibhav Gogate (UW+UCI)	Vibhav Gogate (UW+UCI)	Thomas Schiex (INRA)
3600	Vibhav Gogate (UW+UCI)	Vibhav Gogate (UW+UCI)	Joris Mooij (Max Planck)

toulbar2 was also first at UAI'08 Evaluation, MaxCSP'06/'08 Competition

Winning Teams

- (MAR) **IJGP** by Vibahv Gogate (UW), Andrew Gefland, Natasha Flerova and Rina Dechter (UCI):
Anytime iterative GBP based algorithm
- (PR) **Vgogate** by Vibahv Gogate, Pedro Domingos (UW),, Andrew Gefland and Rina Dechter (UCI):
Formula based importance sampling
- (PR+MAR) **EDBP** by Arthur Choi, Adnan Darwiche, with support from Glen Lenker and Knot Pipatsrisawat (UCLA):
Anytime BP based anytime thickening of structure
- (MAP) **libDAI** by Joris Mooij (Max Planck):
junction tree, LBP/MP, double-loop GBP, Gibbs, decimation
- (MAP) **toulbar2** by Thomas Schiex et al (INRA)
Anytime branch and bound weighted CSP solver

73% instances solved exactly

Toulbar2 & aolib & benchmarks



toulbar2

<http://mulcyber.toulouse.inra.fr/projects/toulbar2>

(Open source WCSP, MPE solver in C++ and counting/PR)



aolib

<http://graphmod.ics.uci.edu/group/Software>

(WCSP, MPE, ILP solver in C++, inference and counting)



Large set of benchmarks

<http://carlit.toulouse.inra.fr/cgi-bin/awki.cgi/SoftCSP>

<http://graphmod.ics.uci.edu/group/Repository>

<http://www.cril.univ-artois.fr/~lecoutre/benchmarks.html>