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### Enumerating Isolated Cliques in Synthetic and Financial Networks

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Task: Enumerate all maximal cliques, that is, all vertex subsets  $C \subseteq V$  such that G[C] is complete and there is no  $C' \supset C$  such that G[C'] is complete.



#### Applications

- ► Computational finance [BOGINSKI ET AL., Comput. Oper. Res., 2006]
- ▶ Biological networks [CHESLER ET AL., Nature Genetics, 2005]
- Social networks, clustering in data mining [MAKINO & UNO, SWAT 2004]

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Clique Enum	neration		

#### Maximal Clique Enumeration

- Simple model
- ▶ NP-hard [Garey & Johnson, 1979]
- ▶ up to 3<sup>n/3</sup> cliques [MOON & MOSER, Israel J. Math., 1965]

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Clique Enum	neration		

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#### Isolated Cliques

- More specific model
- More efficient enumeration algorithms

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c-lsolation			

Definition [ITO, IWAMA, OSUMI, ESA 2005]

A vertex set  ${\it S}$  is called avg-c-isolated if on average the vertices

in S have less than c neighbors outside of S.

Example: avg-2-isolation



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Running times for enumeration of maximal *c*-isolated cliques min-*c*-isolation  $O(2^c \cdot cm + nm)$ avg-*c*-isolation  $O(4^c \cdot c^4m)$ max-*c*-isolation  $O(2.44^c \cdot cm)$ 

[Komusiewicz et al., COCOON 2007]



Known avg-c-isolation algorithm:

- Algorithm enumerates  $O(2^c)$  cliques
- ► Filter out nonmaximal cliques by pairwise comparison → O(4<sup>c</sup>)

#### Idea

Determine maximality independently for each clique.



▶  $S \subseteq D$  with  $D := (\bigcap_{v \in C} N(v)) \setminus C$ 







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- Remaining task: is there a clique S ⊆ D' such that C ∪ S is an avg-c-isolated clique?







#### Maximality Test

- ▶  $S \subseteq D$  with  $D := (\bigcap_{v \in C} N(v)) \setminus C$
- ▶ |D| < c
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- Remaining task: is there a clique S ⊆ D' such that C ∪ S is an avg-c-isolated clique?
- Remove vertices from D' in order of highest degree





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## Theorem

All maximal c-isolated cliques can be enumerated in  $O(2.89^c \cdot c^2 \cdot m)$  time.

 $G_{n,m,p}$  Model [BEHRISCH & TARAZ, Theoret. Comput. Sci., 2006]

Each of n vertices draws each of m features with probability p, and two vertices are connected by an edge iff they have at least one feature in common.

 $\begin{array}{cccc}
 & \bigcirc & \bigcirc & \bigcirc \\
 m = 3 & & & \\
 n = 6 & & & \bigcirc & \bigcirc & \bigcirc \\
\end{array}$ 

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Synthetic I	Data: G <sub>nmn</sub>	Model	

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Each of n vertices draws each of m features with probability p, and two vertices are connected by an edge iff they have at least one feature in common.

	{1} O	{1,2} O	{3} O
<i>m</i> = 3			
<i>n</i> = 6	0	0	0
	$\{1\}$	$\{1, 2\}$	$\{2, 3\}$

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Synthetic D	ata: $G_{n,m,n}$	Model	

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- Each feature induces a clique.
- ► Every nonempty intersection of feature cliques is a clique ~→ we obtain many cliques.

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$\overline{G_{n,m,p}}$ Mo	del		



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G <sub>nmn</sub> Mod	el		



"bk" is an improved variant of the standard Bron-Kerbosch algorithm, which enumerates all maximal cliques.

[KOCH, Theoret. Comput. Sci., 2001]

 $G_{n,m,p}$ 





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Stock Data			

Market Graph [MANTEGNA & STANLEY, 2000]

- Stocks are vertices.
- Two stocks are connected iff the correlation of the daily fluctuations of their prices exceeds some threshold.



(chart data from 2008 Yahoo! Inc.)

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Stock Data			



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Summary			

- Enumerating min- and max-isolated cliques is feasible over a very large range of instances and parameters.
- Sometimes even beats Bron–Kerbosch for enumerating all maximal cliques.
- Avg-isolation more limited.
- Isolation leads to "interesting" cliques, like, e.g., sets of stocks with unusual performance.

#### Other Applications

- ► Finding complexes in protein interaction networks.
- Finding communities in web graphs.
- Finding genres in music artist similarity networks.