

# An Improved Fast Gauss Transform with Applications

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## Fast Multipole Methods

- Originally proposed by Rokhlin and Greengard (1987) to efficiently evaluate sums of monopoles:

- FMM accelerates matrix vector products (sums) of the type

$$\mathbf{v} = \Phi \mathbf{u},$$

- X source point set
- Y evaluation point set
- $\Phi$  some function

- Original functions  $\Phi$  for which FMM was developed were long-ranged and singular at the source point
- FMM relies on “separation of variables” to achieve speed

$$\Phi = \begin{pmatrix} \Phi(\mathbf{y}_1, \mathbf{x}_1) & \Phi(\mathbf{y}_1, \mathbf{x}_2) & \dots & \Phi(\mathbf{y}_1, \mathbf{x}_N) \\ \Phi(\mathbf{y}_2, \mathbf{x}_1) & \Phi(\mathbf{y}_2, \mathbf{x}_2) & \dots & \Phi(\mathbf{y}_2, \mathbf{x}_N) \\ \dots & \dots & \dots & \dots \\ \Phi(\mathbf{y}_M, \mathbf{x}_1) & \Phi(\mathbf{y}_M, \mathbf{x}_2) & \dots & \Phi(\mathbf{y}_M, \mathbf{x}_N) \end{pmatrix}.$$

$$\mathbf{X} = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N\}, \quad \mathbf{x}_i \in \mathbb{R}^d, \quad i = 1, \dots, N,$$

$$\mathbf{Y} = \{\mathbf{y}_1, \mathbf{y}_2, \dots, \mathbf{y}_M\}, \quad \mathbf{y}_j \in \mathbb{R}^d, \quad j = 1, \dots, M.$$

$$\mathbf{v}_j = \sum_{i=1}^N u_i \Phi(\mathbf{y}_j, \mathbf{x}_i), \quad j = 1, \dots, M.$$

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# Factorization

$$\forall \mathbf{x}_i, \mathbf{y}_j \in \Omega \subset \mathbb{R}^d :$$

$$\Phi(\mathbf{y}_j, \mathbf{x}_i) = \sum_{m=0}^{\infty} a_m(\mathbf{x}_i - \mathbf{x}_*) f_m(\mathbf{y}_j - \mathbf{x}_*) = \sum_{m=0}^{p-1} a_m(\mathbf{x}_i - \mathbf{x}_*) f_m(\mathbf{y}_j - \mathbf{x}_*) + Error(p, \mathbf{x}_i, \mathbf{y}_j)$$

Expansion center
Truncation number

Expansion coefficients
Basis functions

- Substitute in the product
- Rearrange summation order
- Inner sum does not depend on evaluation points

$$v_j = \sum_{i=1}^N \Phi(\mathbf{y}_j, \mathbf{x}_i) u_i$$

$$= \sum_{i=1}^N \left[ \sum_{m=0}^{p-1} a_m(\mathbf{x}_i - \mathbf{x}_*) f_m(\mathbf{y}_j - \mathbf{x}_*) + Error(p, \mathbf{x}_i, \mathbf{y}_j) \right] u_i$$

$$= \sum_{m=0}^{p-1} f_m(\mathbf{y}_j - \mathbf{x}_*) \sum_{i=1}^N a_m(\mathbf{x}_i - \mathbf{x}_*) u_i + \sum_{i=1}^N Error(p, \mathbf{x}_i, \mathbf{y}_j) u_i$$

$$= \sum_{m=0}^{p-1} c_m f_m(\mathbf{y}_j - \mathbf{x}_*) + Error(N, p),$$

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## Reduction of Complexity

Straightforward (nested loops):

```

for j = 1, ..., M
  v_j = 0;
  for i = 1, ..., N
    v_j = v_j + Phi(y_j, x_i) u_i;
  end;
end;
    
```

Complexity:  $O(MN)$

If  $p \ll \min(M, N)$  then complexity reduces!

- Remark:  $O(N)$  for fixed  $p$ .
- However, error grows with  $N$
- For fixed error have to increase  $p$  with  $N$

Factorized:

```

for m = 0, ..., p-1
  c_m = 0;
  for i = 1, ..., N
    c_m = c_m + a_m(x_i - x_*) u_i;
  end;
end;
    
```

```

for j = 1, ..., M
  v_j = 0;
  for m = 0, ..., p-1
    v_j = v_j + c_m f_m(y_j - x_*);
  end;
end;
    
```

Complexity:  $O(pN + pM)$

- For geometrically convergent series this introduces a factor of  $\log N$

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# Conventional FMM

- Function  $\Phi$  is singular and a uniformly valid factorization is not available
- Construct patchwork-quilt of overlapping approximations
  - ❑ Local and Multipole Expansions
- Partition sum into a piece that is computed directly and piece that uses factorization.
- Tree data-structures used to reduce the cost of the piece that must be computed directly to that computed via factorization
  - ❑ Translation operators convert one representation to another
- Achieve  $O(N)$  complexity for fixed  $p$
- Remarks
  - ❑ Building data structures is  $O(N \log N)$
  - ❑ For fixed error  $p$  could depend on  $N$  and make complexity  $O(N \log N)$

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# Fast Gauss Transform

- FMM was applied to evaluate sums of Gaussians by Greengard & Strain (1989, 1991)

$$G(y_j) = \sum_{i=1}^N q_i e^{-\|y_j - x_i\|^2 / h^2}, \quad j = 1, \dots, M.$$

↑ Targets      ↑ Sources

$$\begin{bmatrix} G(y_1) \\ G(y_2) \\ \vdots \\ G(y_M) \end{bmatrix} = \begin{bmatrix} e^{-\|x_1 - y_1\|^2 / h^2} & e^{-\|x_2 - y_1\|^2 / h^2} & \dots & e^{-\|x_N - y_1\|^2 / h^2} \\ e^{-\|x_1 - y_2\|^2 / h^2} & e^{-\|x_2 - y_2\|^2 / h^2} & \dots & e^{-\|x_N - y_2\|^2 / h^2} \\ \vdots & \vdots & \ddots & \vdots \\ e^{-\|x_1 - y_M\|^2 / h^2} & e^{-\|x_2 - y_M\|^2 / h^2} & \dots & e^{-\|x_N - y_M\|^2 / h^2} \end{bmatrix} \begin{bmatrix} q_1 \\ q_2 \\ \vdots \\ q_N \end{bmatrix}$$

- Direct evaluation requires  $O(N^2)$  operations.
- FGT reduces cost to  $O(N \log N)$  operations.

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# Original FGT Factorization : Hermite Expansion

- Gaussian kernel factorized into Hermite and Taylor expansions

$$e^{-\|y-x_i\|^2/h^2} = \sum_{n=0}^{p-1} \frac{1}{n!} \left(\frac{x_i - x_*}{h}\right)^n h_n\left(\frac{y - x_*}{h}\right) + \epsilon(p),$$

$$e^{-\|y-x_i\|^2/h^2} = \sum_{n=0}^{p-1} \frac{1}{n!} \left(\frac{y - x_*}{h}\right)^n h_n\left(\frac{x_i - x_*}{h}\right) + \epsilon(p),$$

- where Hermite function  $h_n(x)$  is defined by

$$h_n(x) = (-1)^n \frac{d^n}{dx^n} \left( e^{-x^2} \right).$$

- Exchange order of summations

$$\begin{aligned} G(y_j) &= \sum_{i=1}^N q_i \sum_{n=0}^{p-1} \frac{1}{n!} \left(\frac{x_i - x_*}{h}\right)^n h_n\left(\frac{y_j - x_*}{h}\right) + \epsilon(p), \\ &= \sum_{n=1}^{p-1} A_n h_n\left(\frac{y_j - x_*}{h}\right) + \epsilon(p) \end{aligned}$$

- where  $A_n$  is defined by

$$A_n = \frac{1}{n!} \sum_{i=1}^N q_i \left(\frac{x_i - x_*}{h}\right)^n$$

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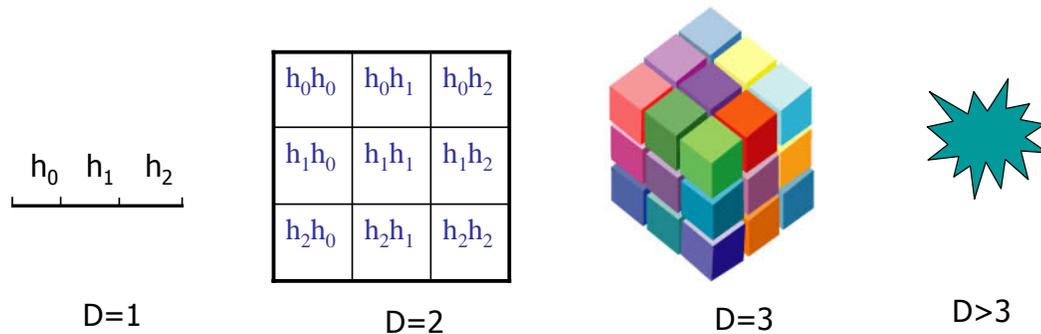
## FGT obtained by applying FMM framework

- Local and “far-field” expansion
- Translation of Hermite expansion to Taylor expansion
- Box data-structures
- Our goal to use the FGT for problems in computer vision and pattern recognition
- Problems not restricted to 1-3 dimensions
  - High dimensional “feature” spaces
- Need to use FGT in high dimensions
- FGT does not scale well with dimensionality

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# Hermite Expansion in Higher Dimensions

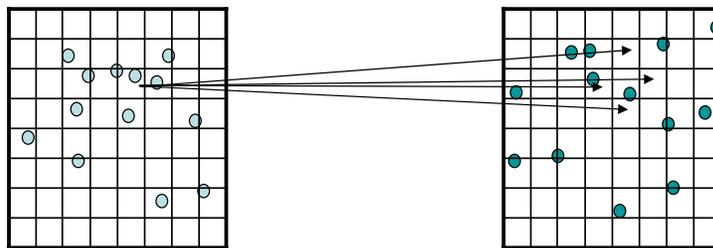
- The higher dimensional Hermite expansion is the Kronecker product of  $d$  univariate Hermite expansions.
- Total number of terms is  $O(p^d)$ ,  $p$  is the number of truncation terms.
- The number of operations in one factorization is  $O(p^d)$ .



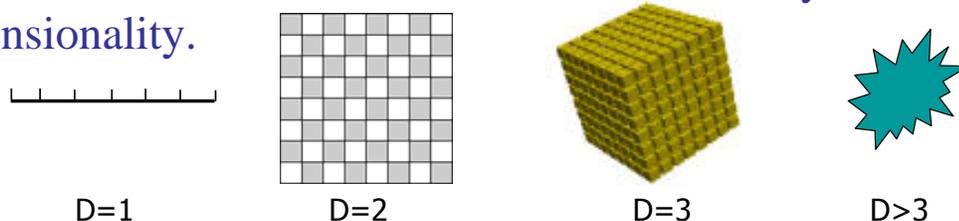
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# Space Subdivision in FGT

- The FGT subdivides the space into uniform boxes and assigns the source points and target points into boxes.
- For each box the FGT maintain a neighbor list.



- The number of the boxes increases exponentially with the dimensionality.



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## FGT in Higher Dimensions

- The higher dimensional Hermite expansion is the product of univariate Hermite expansion along each dimension. Total number of terms is  $O(p^d)$ .
- The space subdivision scheme in the original FGT is uniform boxes. The number of boxes grows exponentially with dimension. Most boxes are empty.
- The FGT was originally designed to solve the problems in mathematical physics (heat equation, vortex methods, etc), where the dimension is up to 3.
- The exponential dependence on the dimension makes the FGT extremely inefficient in higher dimensions.

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## Improved Fast Gauss Transform

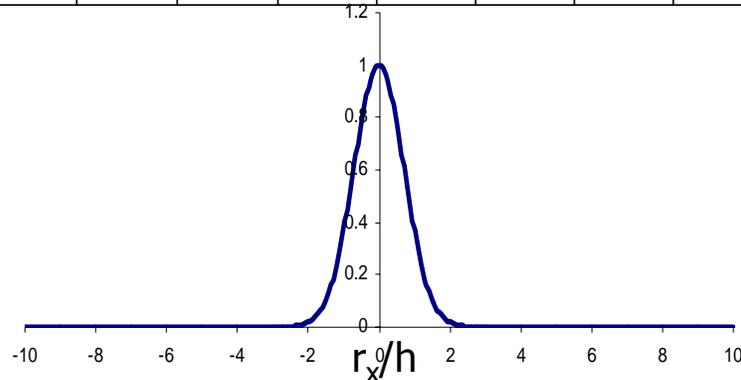
- Reconsider data structures and expansions needed
- Comparing Gaussians with conventional FMM  $\Phi$ 
  - ❑ Gaussian is not singular – it is infinitely differentiable!
  - ❑ Gaussians vanish exponentially quickly in the far-field
- Modified expansions
  - ❑ Local: Multivariate Taylor Expansions
  - ❑ Far field expansion is zero!
- Modified data structures
  - ❑ Data structures are not needed to separate domains of validity (expansions are valid throughout)
  - ❑ Rather need data structures to decide where to ignore the effect of the Gaussian and to decide center of Gaussian

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## Far Field Expansion is Zero

- The decay of the Gaussian kernel function is rapid.
  - Effect of Gaussian outside certain range can be safely ignored
- Time consuming translation operators in original FGT can be safely removed!

$r_x/h$	0	1	2	3	4	5	6	7	8	9	10
$f(r_x/h)$	1	0.367879	0.018316	0.000123	1.13E-07	1.39E-11	2.32E-16	5.24E-22	1.6E-28	6.64E-36	3.72E-44



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## Multivariate Taylor Expansions

- The Taylor expansion of the Gaussian function:

$$e^{-\|y_j - x_i\|^2/h^2} = e^{-\|y_j - x_*\|^2/h^2} e^{-\|x_i - x_*\|^2/h^2} e^{2(y_j - x_*) \cdot (x_i - x_*)/h^2},$$

- The first two terms depend on  $x_i$  or  $y_j$  alone.
- The Taylor expansion of the last term is:

$$e^{2(y_j - x_*) \cdot (x_i - x_*)/h^2} = \sum_{\alpha \geq 0} \frac{2^{|\alpha|}}{\alpha!} \left(\frac{x_i - x_*}{h}\right)^\alpha \left(\frac{y_j - x_*}{h}\right)^\alpha.$$

where  $\alpha = (\alpha_1, \dots, \alpha_d)$  is multi-index.

- The multivariate Taylor expansion about center  $x_*$ :

$$G(y_j) = \sum_{\alpha \geq 0} C_\alpha e^{-\|y_j - x_*\|^2/h^2} \left(\frac{y_j - x_*}{h}\right)^\alpha,$$

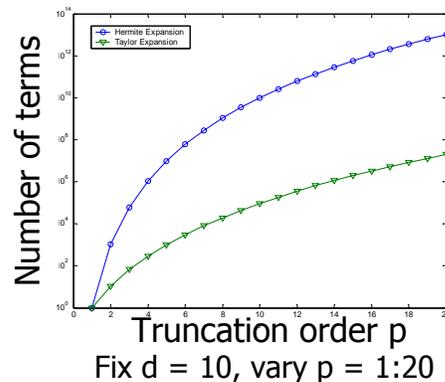
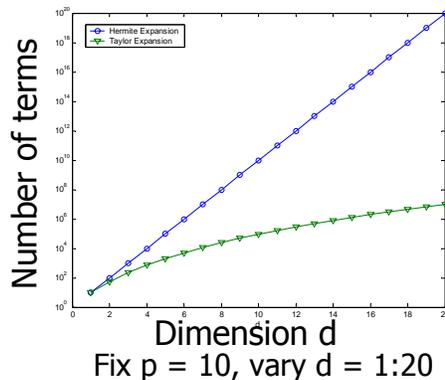
- where coefficients  $C_\alpha$  are given by

$$C_\alpha = \frac{2^{|\alpha|}}{\alpha!} \sum_{i=1}^N q_i e^{-\|x_i - x_*\|^2/h^2} \left(\frac{x_i - x_*}{h}\right)^\alpha.$$

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# Modified Factorization: Taylor Expansions

- The number of terms in multivariate Taylor expansion is  $\binom{p+d-1}{d}$ , asymptotically  $O(d^p)$
- Original expansion has  $O(p^d)$  terms
- New expansion results in a big reduction for large  $d$  and moderate  $p$



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## Space Subdivision Scheme

- The space subdivision scheme in the original FGT is uniform boxes. The number of boxes grows exponentially with the dimensionality.
- Need a data structure that
  - ❑ Allows ignoring the far-field
  - ❑ Assigns each point to a local expansion center
- The space subdivision should adaptively fit density of the points.
- The cell should be as compact as possible.
- The algorithm should be a progressive one,
  - ❑ Refined space subdivision obtained from previous one.
- Based on the above considerations, we develop a structure using the  $k$ -center problem.

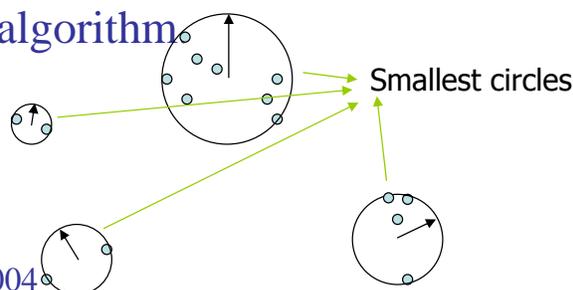
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# $k$ -center Algorithm

- The  $k$ -center problem is defined to seek the “best” partition of a set of points into clusters (Gonzalez 1985, Hochbaum and Shmoys 1985, Feder and Greene 1988).

Given a set of points and a predefined number  $k$ ,  $k$ -center clustering is to find a partition  $S = S_1 \cup S_2 \cup \dots \cup S_k$  that minimizes  $\max_{1 \leq i \leq k} \text{radius}(S_i)$ , where  $\text{radius}(S_i)$  is the radius of the smallest disk that covers all points in  $S_i$ .

- The  $k$ -center problem is NP-hard but there exists a simple 2-approximation algorithm.



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# Farthest-Point Algorithm

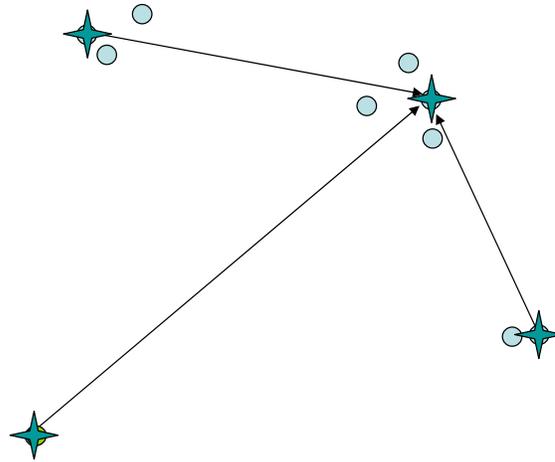
- The farthest-point algorithm (a.k.a.  $k$ -center algorithm) is a 2-approximation to optimal solution (Gonzales 1985).
- The total running time is  $O(kn)$ ,  $n$  is the number of points. It can be reduced to  $O(n \log k)$  using a slightly more complicated algorithm (Feder and Greene 1988).

- Initially randomly pick a point  $v_0$  as the first center and add it to the center set  $C$ .
- For  $i = 1$  to  $k - 1$  do
  - For every point  $v \in V$ , compute the distance from  $v$  to the current center set  $C = \{v_0, v_1, \dots, v_{i-1}\}$ :  $d_i(v, C) = \min_{c \in C} \|v - c\|$ .
  - From the points  $V - C$  find a point  $v_i$  that is farthest away from the current center set  $C$ , i.e.  $d_i(v_i, C) = \max_v \min_{c \in C} \|v - c\|$ .
  - Add  $v_i$  to the center set  $C$ .
- Return the center set  $C = \{v_0, v_1, \dots, v_{k-1}\}$  as the solution to  $k$ -center problem.

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# A Demo of $k$ -center Algorithm

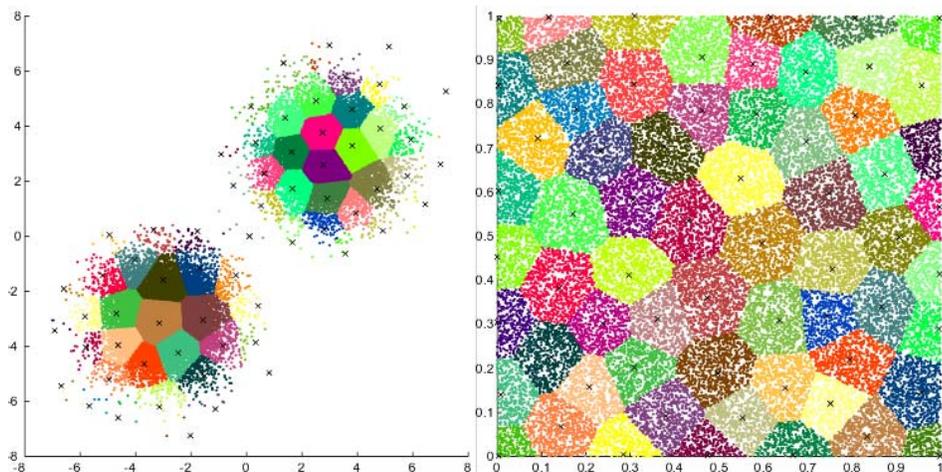
$k = 4$



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## Results of $k$ -center Algorithm

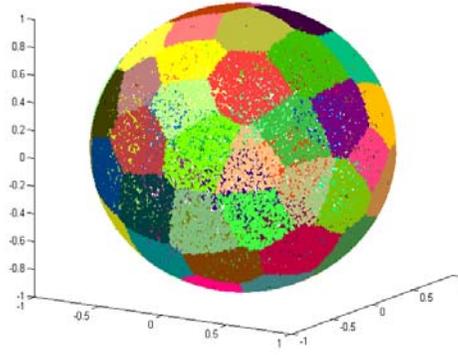
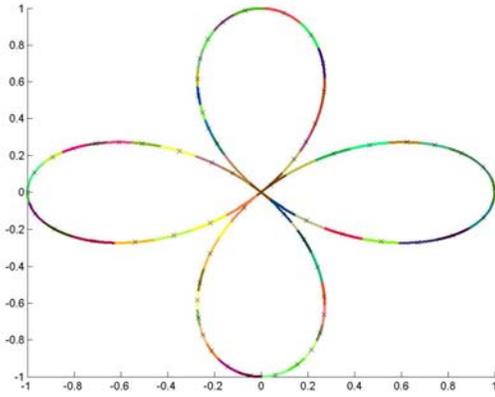
- The results of  $k$ -center algorithm. 40,000 points are divided into 64 clusters in 0.48 sec on a 900MHZ PIII PC.



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# More Results of k-center Algorithm

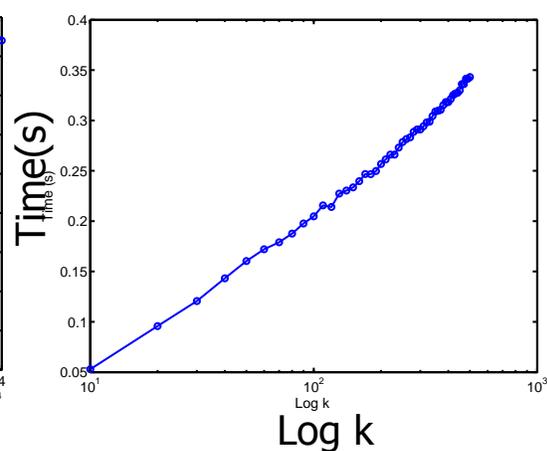
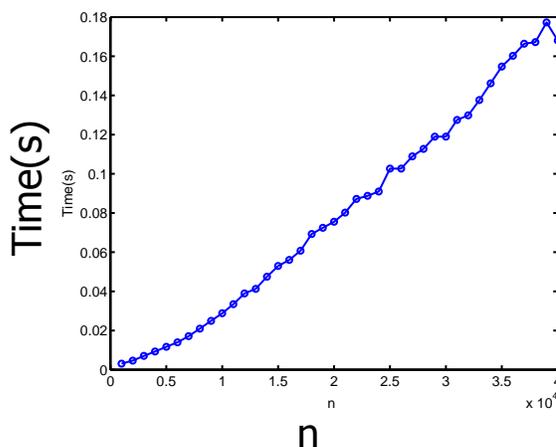
- The 40,000 points are on the manifolds.



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## Properties of $k$ -center Algorithm

- Computational complexity of  $k$ -center is  $O(n \log k)$ .
  - Points are generated using uniform distribution.
  - (Left) Number of points varies from 1000 to 40000 for  $k=64$
  - (Right) Number of clusters  $k$  varies from 10 to 500 for 40000 points.



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# Monomial Orders

- Let  $\alpha=(\alpha_1, \dots, \alpha_n), \beta=(\beta_1, \dots, \beta_n)$ , then three standard monomial orders:
  - Lexicographic order, or “dictionary” order:**
    - $\alpha \prec_{\text{lex}} \beta$  iff the leftmost nonzero entry in  $\alpha - \beta$  is negative.
  - Graded lexicographic order:**
    - $\alpha \prec_{\text{grlex}} \beta$  iff  $\sum_{1 \leq i \leq n} \alpha_i < \sum_{1 \leq i \leq n} \beta_i$  or  $(\sum_{1 \leq i \leq n} \alpha_i = \sum_{1 \leq i \leq n} \beta_i$  and  $\alpha \prec_{\text{lex}} \beta$ ).
  - Graded reverse lexicographic order:**
    - $\alpha \prec_{\text{grevlex}} \beta$  iff  $\sum_{1 \leq i \leq n} \alpha_i < \sum_{1 \leq i \leq n} \beta_i$  or  $(\sum_{1 \leq i \leq n} \alpha_i = \sum_{1 \leq i \leq n} \beta_i$  and the rightmost nonzero entry in  $\alpha - \beta$  is positive).
- Example:**
  - Let  $f(x,y,z) = xy^5z^2 + x^2y^3z^3 + x^3$ , then
    - w.r.t. lex:  $f(x,y,z) = x^3 + x^2y^3z^3 + xy^5z^2$ ;
    - w.r.t. grlex:  $f(x,y,z) = x^2y^3z^3 + xy^5z^2 + x^3$ ;
    - w.r.t. grevlex:  $f(x,y,z) = xy^5z^2 + x^2y^3z^3 + x^3$ .

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# Horner's Rule

- Horner's rule (Horner, 1819) *recursively* evaluates the polynomial  $a_p x^p + \dots + a_1 x + a_0$  as:
 
$$((\dots(a_p x + a_{p-1})x + \dots)x + a_0.$$
- costs  $p$  multiplications and  $p$  additions, no extra storage.
  - Reduces complexity from  $O(p^2)$  to  $O(p)$
- We do this for the multivariate polynomial *iteratively* using the graded lexicographic order. Costs  $C(p+d-1, d)$  operations and storage.

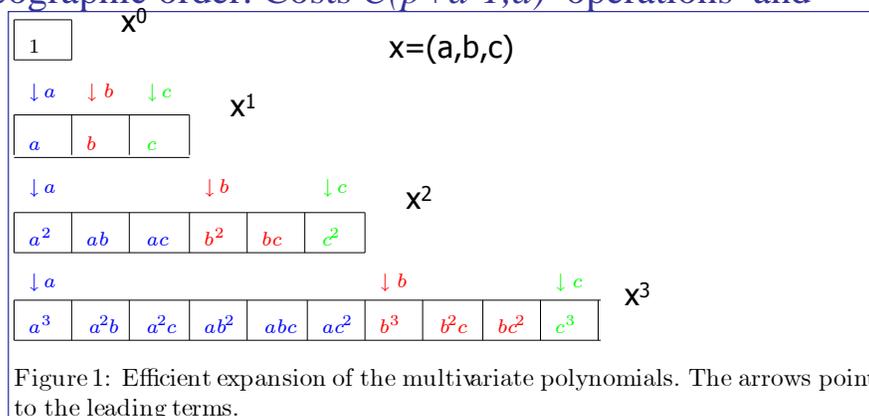


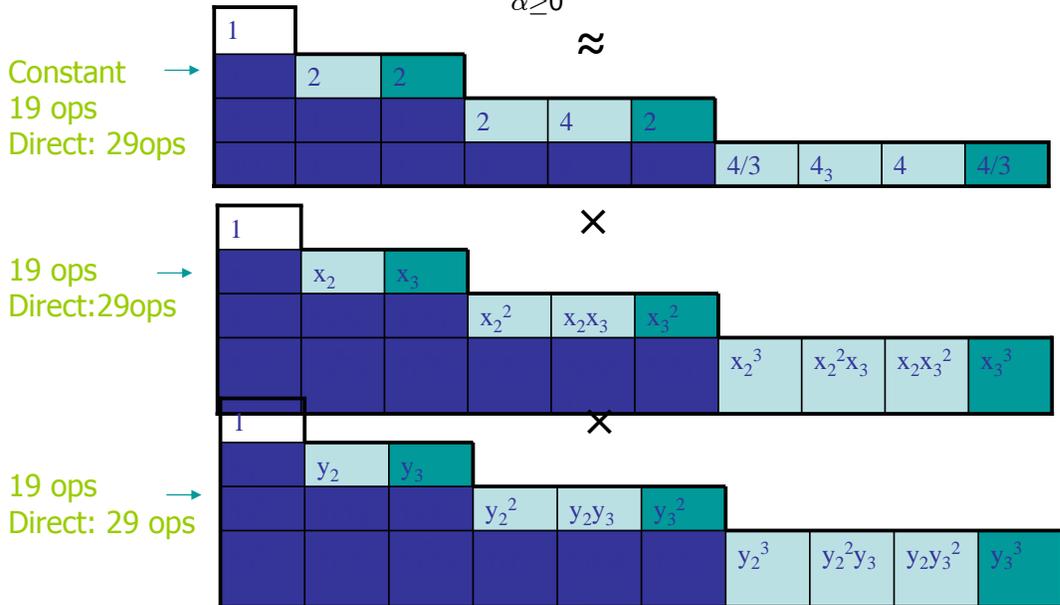
Figure 1: Efficient expansion of the multivariate polynomials. The arrows point to the leading terms.

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# An Example of Taylor Expansion

- Suppose  $x = (x_1, x_2, x_3)$  and  $y = (y_1, y_2, y_3)$ , then

$$e^{2x \cdot y} = \sum_{\alpha \geq 0} \frac{2^{|\alpha|}}{\alpha!} x^\alpha y^\alpha \approx$$

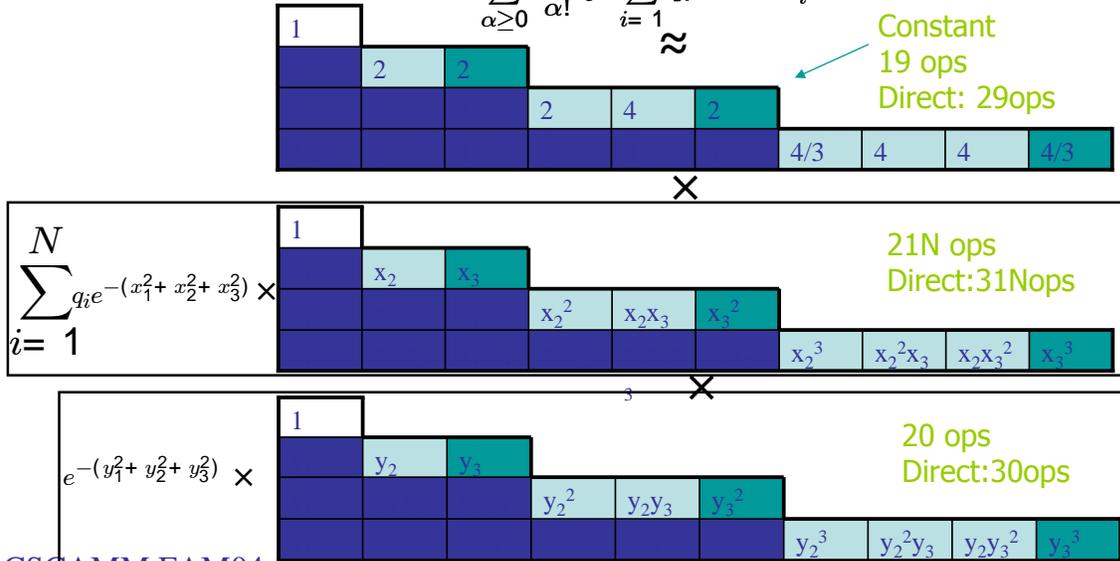


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# An Example of Taylor Expansion (Cont'd)

$$G(y) = \sum_{i=1}^N q_i e^{-\|x_i - y\|^2} = \sum_{i=1}^N q_i e^{-\|x_i\|^2} e^{-\|y\|^2} \sum_{\alpha \geq 0} \frac{2^{|\alpha|}}{\alpha!} x_i^\alpha y^\alpha$$

$$= e^{-\|y\|^2} \sum_{\alpha \geq 0} \frac{2^{|\alpha|}}{\alpha!} y^\alpha \sum_{i=1}^N q_i e^{-\|x_i\|^2} x_i^\alpha \approx$$



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# Improved Fast Gauss Transform

Control series truncation error

**Step 1** Assign the  $N$  sources into  $K$  clusters using the farthest-point clustering algorithm such that the radius is less than  $r_x$ .

**Step 2** Choose  $p$  sufficiently large such that the error estimate is less than the desired precision  $\epsilon$ .

**Step 3** For each cluster  $S_k$  with center  $c_k$ , compute the coefficients:

$$C_\alpha^k = \frac{2^{|\alpha|}}{\alpha!} \sum_{x_i \in S_k} q_i e^{-\|x_i - c_k\|^2/h^2} \left( \frac{x_i - c_k}{h} \right)^\alpha.$$

Collect the contributions from sources to centers

**Step 4** Repeat for each target  $y_j$ , find its neighbor clusters whose centers lie within the range  $r_y$ . Then the sum of Gaussians can be evaluated by the expression:

$$G(y_j) = \sum_{\|y_j - c_k\| < h r_y} \sum_{|\alpha| < p} C_\alpha^k e^{-\|y_j - c_k\|^2/h^2} \left( \frac{y_j - c_k}{h} \right)^\alpha.$$

Control far field cutoff error

Summarize the contributions from centers to targets

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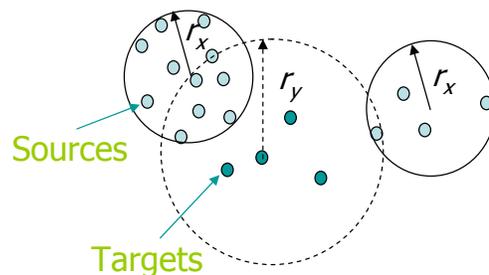
## Error Bound of IFGT

- The total error from the series truncation and the cutoff outside of the neighborhood of targets is bounded by

$$|E(y)| \leq \sum |q_i| \left( \frac{2^p}{p!} \left( \frac{r_x}{h} \right)^p \left( \frac{r_y}{h} \right)^p + e^{-(r_y/h)^2} \right).$$

Truncation error

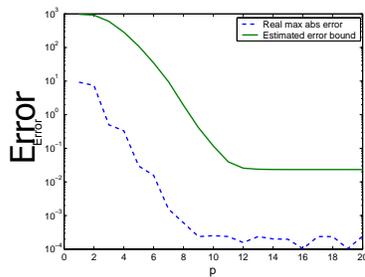
Cutoff error



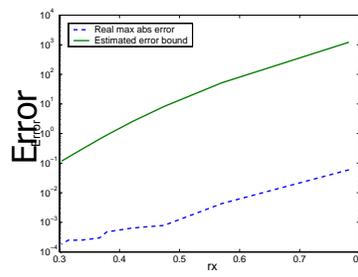
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# Error Bound Analysis

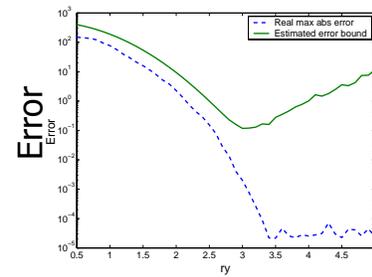
- Increasing number of truncation terms  $p$ , reduces error
- Increasing  $k$  in the  $k$ -center algorithm, radius of source point clusters  $r_x$  will decrease, until the error bound is less than a given precision.
- The error bound first decreases, then increases with respect to the cutoff radius  $r_y$ .



Truncation order  $p$



Max radius of cells

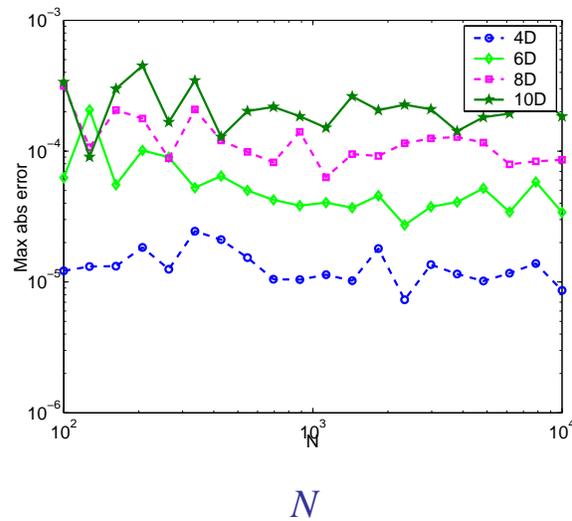
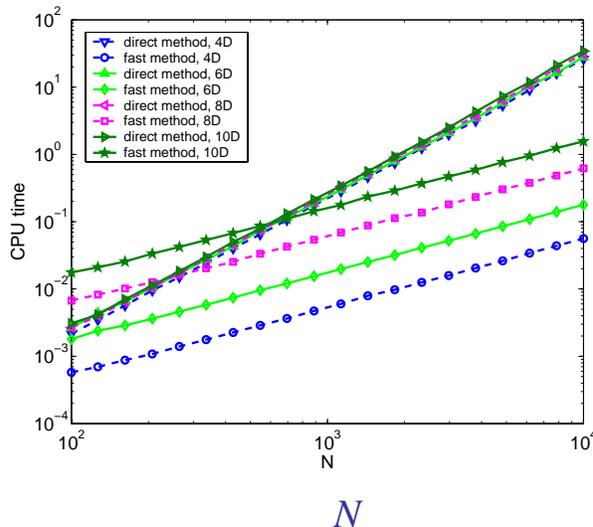


Cutoff radius

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# Experimental Result

- The speedup of the fast Gauss transform in 4, 6, 8, 10 dimensions ( $h=1.0$ ).



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# Efficient Kernel Density Estimation

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## Kernel Density Estimation (KDE)

- Kernel density estimation (a.k.a Parzen method, Rosenblatt 1956, Parzen 1962) is an important nonparametric technique.
- KDE is the keystone of many algorithms:
  - ❑ Radial basis function networks
  - ❑ Support vector machines
  - ❑ Mean shift algorithm
  - ❑ Regularized particle filter
- The main drawback is the quadratic computational complexity. Very slow for large dataset.

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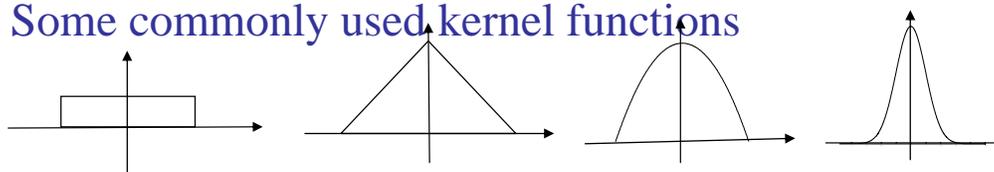
# Kernel Density Estimation

- Given a set of observations  $\{x_1, \dots, x_n\}$ , an estimate of density function is

$$\hat{f}_n(\mathbf{x}) = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{\|\mathbf{x} - \mathbf{x}_i\|}{h}\right)$$

Dimension  $\leftarrow$   $nh^d$       Kernel function  $\leftarrow$   $K\left(\frac{\|\mathbf{x} - \mathbf{x}_i\|}{h}\right)$       Bandwidth  $\leftarrow$   $h$

- Some commonly used kernel functions



- The computational requirement for large datasets is  $O(N^2)$ , for  $N$  points.

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## Efficient KDE and FGT

- In practice, the most widely used kernel is the Gaussian

$$K_N(\mathbf{x}) = (2\pi)^{-d/2} e^{-\frac{1}{2}\|\mathbf{x}\|^2}.$$

- The density estimate using the Gaussian kernel:

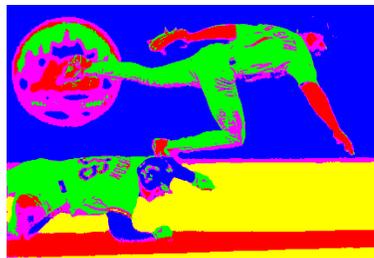
$$\hat{p}_n(\mathbf{x}) = c_N \sum_{i=1}^N e^{-\|\mathbf{x} - \mathbf{x}_i\|^2/h^2}$$

- Fast Gauss transform can reduce the cost to  $O(N \log N)$  in low-dimensional spaces.
- Improved fast Gauss transform accelerates the KDE in both lower and higher dimensions.

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# Experimental Result

- Image segmentation results of the mean-shift algorithm with the Gaussian kernel.



Size: 432X294  
Time: 7.984 s

Direct evaluation:  
more than 2 hours



Size: 481X321  
Time: 12.359 s

Direct evaluation:  
more than 2 hours

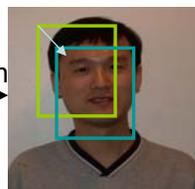
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## Object Tracking

- Goal of object tracking: find the moving objects between consecutive frames.
- A model image or template is given for tracking.
- Usually a feature space is used, such as pixel intensity, colors, edges, etc.
- Usually a similarity measure is used to measure the difference between the model image and current image.
- Temporal correlation assumption: the change between two consecutive frames is small.



Similarity function



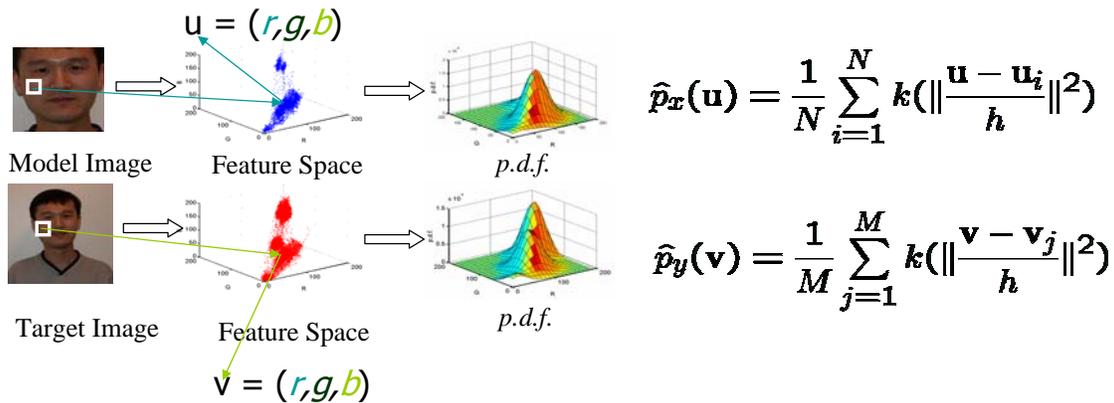
Model image

Target image

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# Image Representations

- Images are mapped into feature spaces.
- Feature spaces are described by the probabilistic density functions (*p.d.f.*).
- The *p.d.f.* is estimated using kernel density estimation:



- Accelerated using FGT. Details in Yang et al 2004.

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## Experimental results



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# Experimental results



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## Future Work

- Applications to classification via dimension reduction and FGT accelerated SVM
- Bandwidth selection
- FGT code FIGTREE (v 1.0) to be released shortly
  - ❑ Free for noncommercial use

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