# **Orthogonal Eigenvectors and Gram-Schmidt**

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- One of FOUR main papers on this work
- Algorithm  $MR^3$  or MRRR
  - Acronym for Multiple Relatively Robust Representations
  - Accurate but turgid title
  - Jim Demmel has a more catchy title...
- Guiding Principle: No Gram-Schmidt

interest very close to 0. The middle part of this paper presents the relevant error analysis. Although essential for our results, this analysis will be indigestible for most readers, but it tells us that changes of only 3 or 4 units in the last digit of each entry of the input L, D and the output twisted factors suffice to give the exact relation.

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  - Stick closely to the paper

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- Beautiful identity:

$$D_{+} + D_{-} = \operatorname{diag}(J) + \operatorname{diag}(J^{-1})^{-1}$$

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  - Analyze with a Representation Tree

# Difficulties

- All eigenvalues of *T* are easily computed in  $O(n^2)$  time
- Given  $\hat{\lambda}$ , inverse iteration computes the eigenvector:

$$(T - \hat{\lambda}I)x_{i+1} = x_i, \quad i = 0, 1, 2, \dots$$

- Costs O(n) per iteration
- Typically, 1-3 iterations are enough
- **BUT**, inverse iteration only guarantees

$$\|T\hat{v} - \hat{\lambda}\hat{v}\| = O(\varepsilon \|T\|)$$

## **Fundamental Limitations**

Gap Theorem :

$$\sin \angle (v, \hat{v}) \le rac{\|T\hat{v} - \hat{\lambda}\hat{v}\|}{\mathsf{Gap}(\hat{\lambda})}$$

 $\operatorname{Gap}(\hat{\lambda})$  can be small :

$$\begin{bmatrix} 1 & \varepsilon_1 \\ \varepsilon_1 & 1 & \varepsilon_2 \\ & \varepsilon_2 & 1 & \varepsilon_3 \\ & & \varepsilon_3 & 1 \end{bmatrix}$$

When eigenvalues are close, independently computed eigenvectors **WILL NOT** be mutually orthogonal

## **Example from Quantum Chemistry**

- Symmetric positive definite eigenproblem, n = 966
- Occurs in Møller-Plesset theory in the modeling of biphenyl
- Eigenvalue Distribution:



• LAPACK "clusters" — numbered  $1, \ldots, 939$  and smaller ones

### **Example from Quantum Chemistry**

Plot of  $Absgap(i) = \log_{10}(\min(\lambda_{i+1} - \lambda_i, \lambda_i - \lambda_{i-1})/||T||)$  versus i:



Plot of  $\operatorname{Relgap}(i) = \log_{10}(\min(\lambda_{i+1} - \lambda_i, \lambda_i - \lambda_{i-1})/|\lambda_i|)$  versus i:



# **First Steps**

- Factor  $T = LDL^T$
- Compute eigenvalues of  $LDL^T$  by dqds or bisection
- For each eigenvalue  $\lambda$ , compute eigenvector by inverse iteration

## **Computing Eigenvector #1**

- $\hat{\lambda}_1 = 4.1338 \times 10^{-08}, \, \hat{\lambda}_2 = 4.3417 \times 10^{-08}, \, \hat{\lambda}_3 = 4.5 \dots \times 10^{-08}$
- Factor  $T_1 = LDL^T \hat{\lambda}_1 I = L_+ D_+ L_+^T = U_- D_- U_-^T$  (up & down)
- Compute  $\gamma(i) = D_{+}(i) + D_{-}(i) T_{1}(i,i)$



• Solve  $T_1 z_1 = \gamma_r e_r$ , where  $\gamma_r = \min_k |\gamma_k|$ 



## **Computing Eigenvector #2**

- $\hat{\lambda}_1 = 4.1338 \times 10^{-08}, \, \hat{\lambda}_2 = 4.3417 \times 10^{-08}, \, \hat{\lambda}_3 = 4.5 \dots \times 10^{-08}$
- Factor  $T_2 = LDL^T \hat{\lambda}_2 I = L_+ D_+ L_+^T = U_- D_- U_-^T$  (up & down)
- Compute  $\gamma(i) = D_+(i) + D_-(i) T_2(i,i)$



• Solve  $T_2 z_2 = \gamma_r e_r$ , where  $\gamma_r = \min_k |\gamma_k|$ 



- $|z_1^T z_2| < 2\varepsilon$
- Residual norms are  $< \varepsilon \|T\|$

In general,

- $\min_k |\gamma_k| \approx \varepsilon |\hat{\lambda}|$
- Dot Product between two vectors  $\propto \frac{\varepsilon |\lambda|}{\text{Gap}}$
- If two eigenvalues share d leading digits, then dot product  $\approx 10^d \varepsilon$

•  $\hat{\lambda}_{280}$  to  $\hat{\lambda}_{303}$  are :

0.2852950617

0.2877656004

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- Some eigenvalues agree in 5 digits
- Compute new shifted representations:

$$\begin{split} LDL^{T} &- 0.2852950617I = L_{1}D_{1}L_{1}^{T} \\ \hline 0.60738244647 \times 10^{-15} \\ \hline 0.24705386756 \times 10^{-02} \\ \vdots \\ \hline 0.4811762683 \times 10^{-02} \\ \hline 0.4852987244 \times 10^{-02} \\ \hline 0.4855354851 \times 10^{-02} \end{split}$$

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## The computed vectors

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### • Maximum Dot Product $< 3\varepsilon$ .

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$$T \longrightarrow LDL^T$$

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- Seminal 1991 Demmel-Kahan paper (SIAG/LA Prize)
  - Small relative changes in the entries of a bidiagonal cause small relative changes in all its singular values
- L and D almost always define the small eigenvalues of  $LDL^T$  to high relative accuracy
  - Even in the face of element growth!

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$$|x^T y| \le \max(\varepsilon_1, \varepsilon_2) / \frac{|\hat{\lambda} - \hat{\mu}|}{|\hat{\lambda}| + |\hat{\mu}|}$$

Thus, orthogonality depends on the relative separation

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- Thus, orthogonality depends on the relative separation
- Relative Gaps can be made larger by shifting

$$LDL^T - \xi I = \tilde{L}\tilde{D}\tilde{L}^T$$

- Different representations for different clusters!
  - Essential to use differential transforms

### **Differential Transforms**

$$LDL^T - \hat{\lambda}I = L_+D_+L_+^T$$

Simple qd :  $D_{+}(1) := d_{1} - \hat{\lambda}$ for i = 1, n - 1  $L_{+}(i) := (d_{i}l_{i})/D_{+}(i)$   $D_{+}(i+1) := d_{i}l_{i}^{2} + d_{i+1} - L_{+}(i)d_{i}l_{i} - \hat{\lambda}$ end for Differential qd :  $s_{1} := -\hat{\lambda}$ 

for 
$$i = 1, n - 1$$
  
 $D_{+}(i) := s_{i} + d_{i}$   
 $L_{+}(i) := (d_{i}l_{i})/D_{+}(i)$   
 $s_{i+1} := L_{+}(i)l_{i}s_{i} - \hat{\lambda}$   
end for

 $D_+(n) := s_n + d_n$ 

## Key Idea 3 — Twist, again with differential transformations

- Godunov et al. [1985], Fernando [1995]
- Compute the appropriate Twisted Factorization :

$$LDL^T - \hat{\lambda}I = N_r D_r N_r^T,$$



• Solve for z,  $N_r D_r N_r^T z = \gamma_r e_r$  (  $\Rightarrow N_r^T z = e_r$  ):

$$z(i) = \begin{cases} 1, & i = r, \\ -L_{+}(i) \cdot z(i+1), & i = r-1, \dots, 1, \\ -U_{-}(i-1) \cdot z(i-1), & i = r+1, \dots, n. \end{cases}$$

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• Eigenvalues:  $\varepsilon$ ,  $1 + \sqrt{\varepsilon}$ ,  $1 + 2\sqrt{\varepsilon}$ , 2

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• Following Representation Tree captures the steps of the algorithm:



# Caveats

- BLAS
  - Complexity is O(nk) to compute k eigenpairs
  - However, all operations are BLAS 1
  - Closest competitor D&C  $O(n^3)$ , but BLAS 3
- Very tight eigenvalue clusters
  - C. Vömel's torture tests

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Roundoff to the rescue:

$$\lambda_{20}(L_0 D_0 L_0^T)$$
 &  $\lambda_{21}(L_0 D_0 L_0^T)$  — no digits in common!  
-7.28 × 10<sup>-14</sup> & -1.22 × 10<sup>-15</sup>

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• Computed Eigenvectors  $\hat{v}_{20}$  and  $\hat{v}_{21}$  (inner product is  $1.0 \times 10^{-16}$ ):



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- Very tight eigenvalue clusters
  - C. Vömel's torture tests 5 copies of  $W_{101}^+$  with glue of  $\sqrt{\varepsilon}$
  - Required a tweak Perturb base representation