

I Don't Care about BLAS

Devin Matthews

Institute for Computational
Engineering and Sciences, UT Austin

#smallmoleculesmatter

From QC to DGEMM

$$\bar{H} \hat{R} |\Phi\rangle = E \hat{R} |\Phi\rangle \quad \text{“Simple” eigenproblem...}$$

$$r_{ijkl}^{abef}, \bar{H}_{cdkl}^{abij} \quad \text{In terms of tensors...}$$

$$r_{ijkl}^{abef}, W_{ck}^{ai}, F_k^i, t_{ij}^{ab}, \dots \quad \text{In terms of other tensors...}$$

$$r_{ijk\bar{l}}^{ab\bar{e}\bar{f}} \text{ or } \check{r}_{ijkl}^{abef} \quad \text{With structured sparsity...}$$

$$r_{i < j \bar{k} < \bar{l}}^{a < b \bar{e} < \bar{f}} \text{ or } \check{r}_{i \leq j \leq k \leq l}^{abef} \quad \text{With symmetry...}$$

$$r_{0000}^{abef}, r_{0001}^{abef}, r_{0002}^{abef}, \dots \quad \text{With slicing (or blocking etc.)...}$$

$$r_{(ef)\gamma_{ef}}^{(ab)\gamma_{ab}} \quad \text{With more sparsity...}$$

$$r_{ef}^{ab} \in \mathbb{R}^{n_a \otimes n_b \otimes n_e \otimes n_f} \quad \text{In terms of dense tensors...}$$

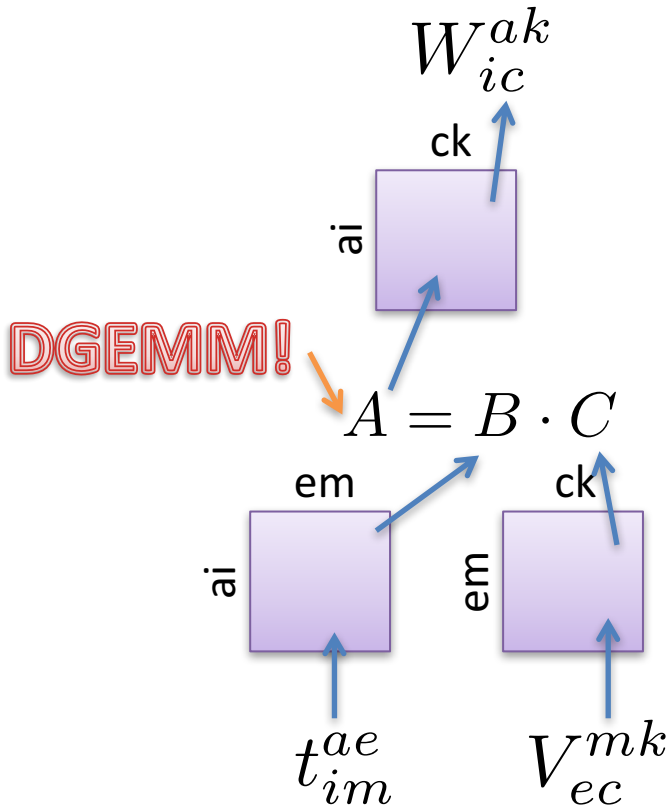
DGEMM!

$$\longrightarrow A = B \cdot C$$

In terms of matrices.

Tensor Contraction Today

“TTDT”



“LoG”

$$W_{ic}^{ak} = t_{im}^{ae} \cdot V_{ec}^{mk}$$

$$[W_{ak}]_{ic} = \sum_m [t_{am}]_{ei} \cdot [V_{mk}]_{ec} \quad \forall i, e, c$$

for i
 for e
 for c

DGEMM! → DGEMM

DGEMM



DGEMM Considered Harmful

- Tensors have to be **transposed** in order to use DGEMM.
- DGEMM needs **dense** matrices. If our tensors have **structure** (permutational symmetry, point group symmetry, sparsity, etc.) we have to **expand** or **block** them.
- Point group symmetry is efficiently handled with the **Direct Product Decomposition** (DPD), but we want to *automate* and *optimize* it.
- **Blocking** reduces the size of individual DGEMM calls. Can we **aggregate** these into more efficient operations?

DPD: Stanton, J.F.; Gauss, J.; Watts, J.D.; Bartlett, R.J. *J. Chem. Phys.* **1991**, *94*, 4334.

How Much Does Transpose Cost?

Speedup of NCC (new code) relative to MRCC:

	HSOH	H ₂ O	H ₂ C ₄ H ₂	O ₃	FO ₃ ⁻
CCSDTQ	6.2	4.4	5.2	6.2	4.9
CCSDT(Q)	33.1	102.6	18.2	28.7	17.2

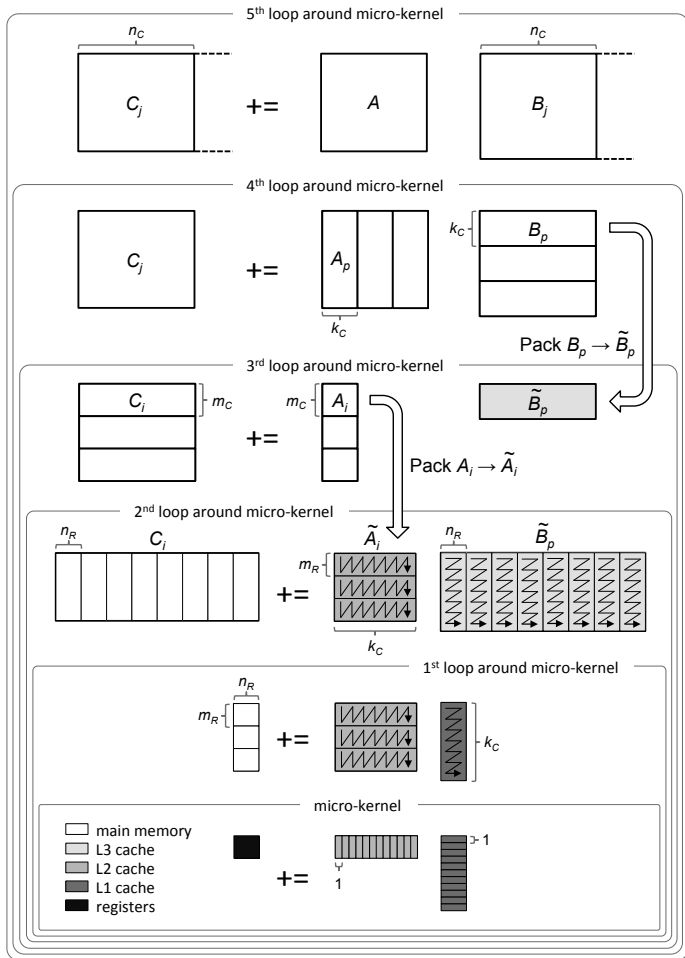
Timing breakdown of (Q) by low-level operation

Level 1 BLAS	2.4%
Level 2 BLAS	2.0%
Level 3 BLAS	47.9%
Disk I/O	< 0.1%
Spin-summation	3.7%
Transpose	41.1%

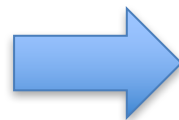
Timing breakdown by low-level operation

Level 1 BLAS	10.9%
Level 2 BLAS	0.9%
Level 3 BLAS	45.5%
Disk I/O	3.4%
Spin-summation	13.0%
Transpose	26.3%

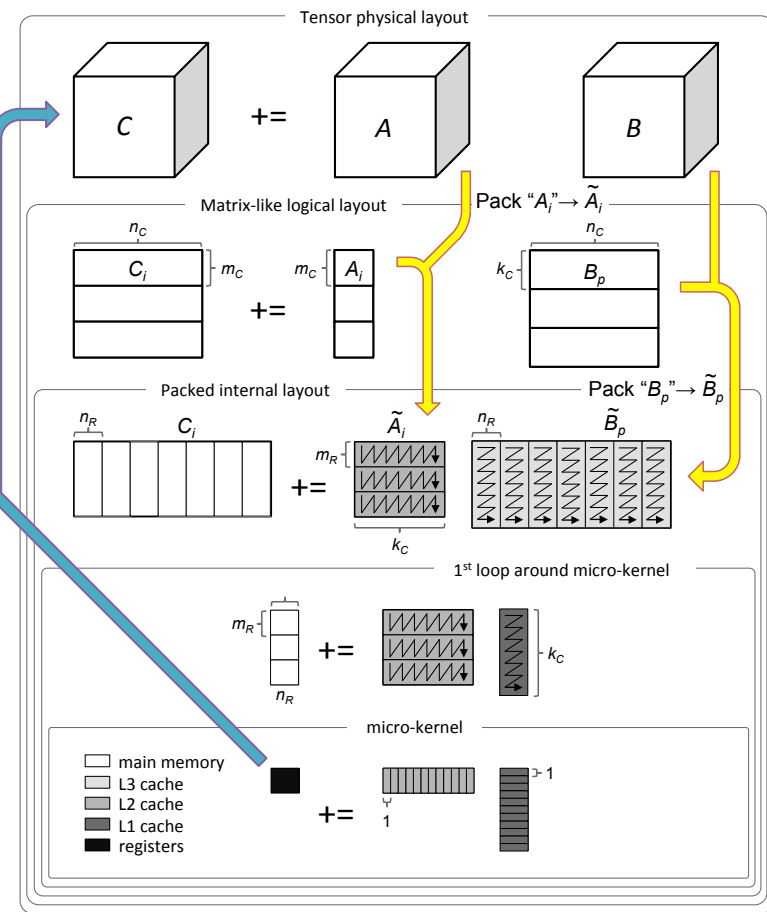
BLIS \rightarrow TBLIS



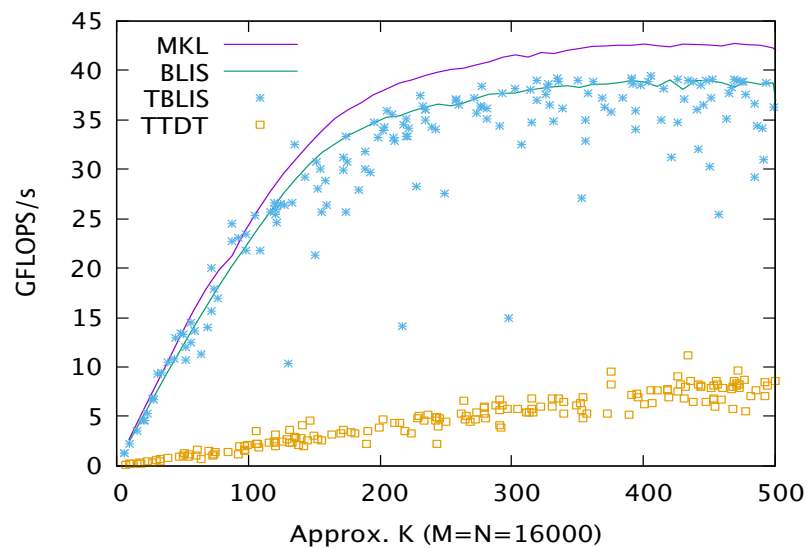
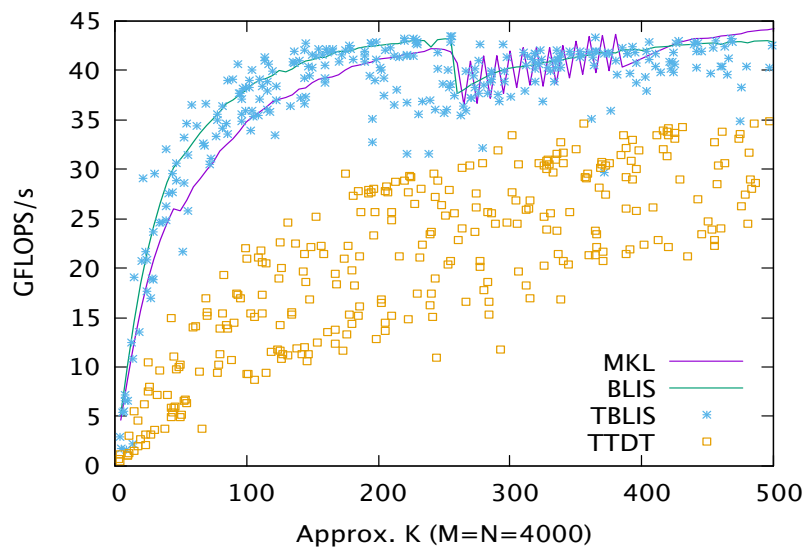
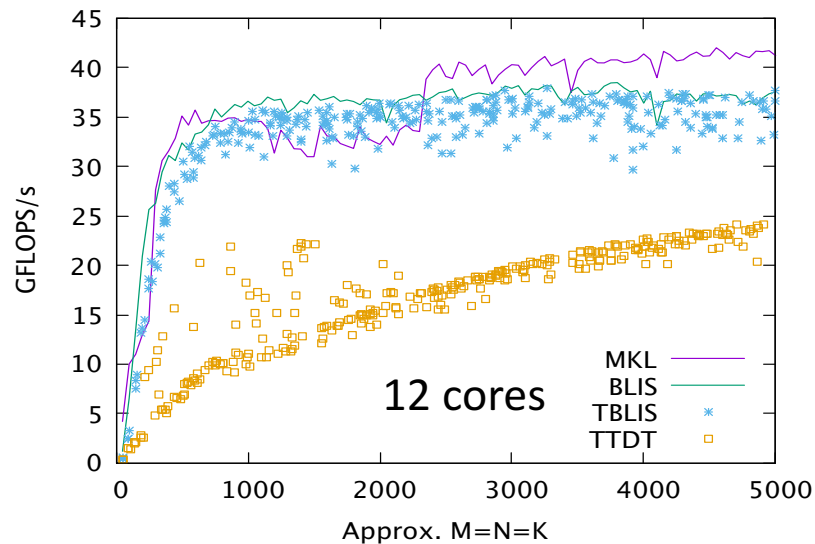
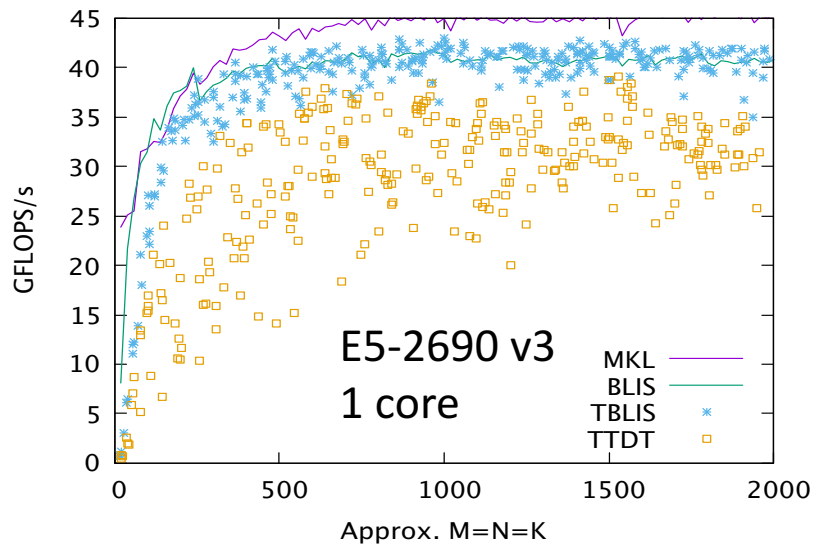
Tensor
to
matrix
packing
kernel



(Possibly)
scattered
update

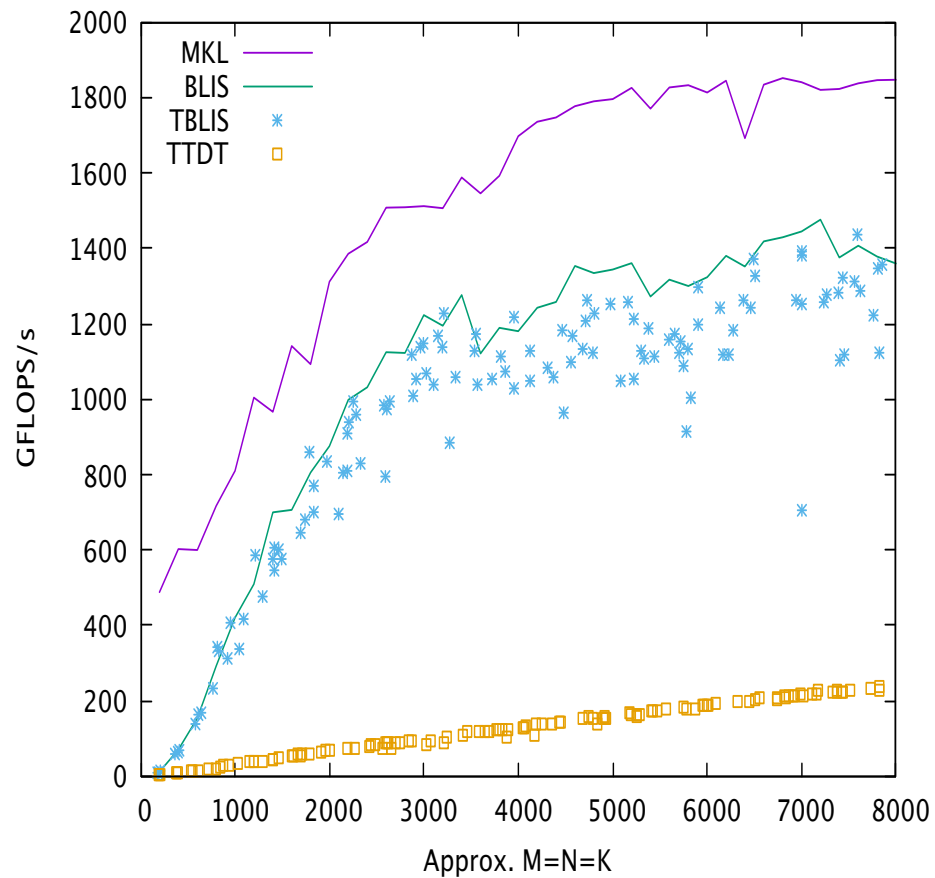


Results for Dense Tensors

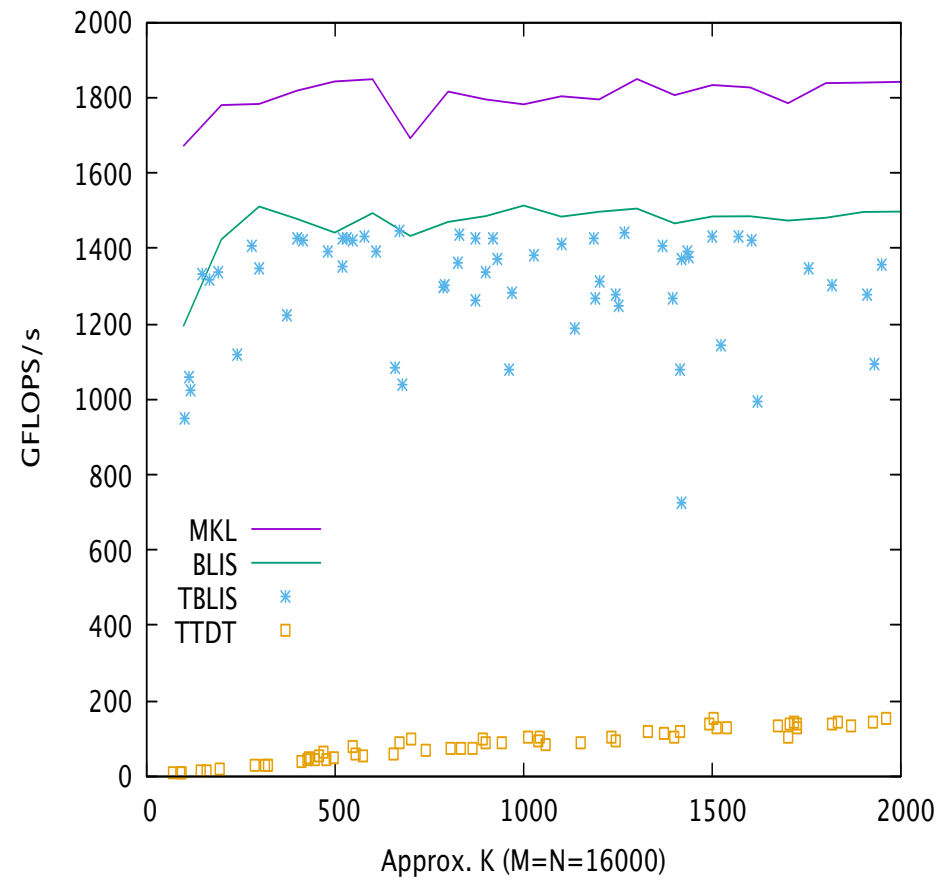


Works Great on Xeon Phi Too

“Square” MM and TC
on Xeon Phi 7210

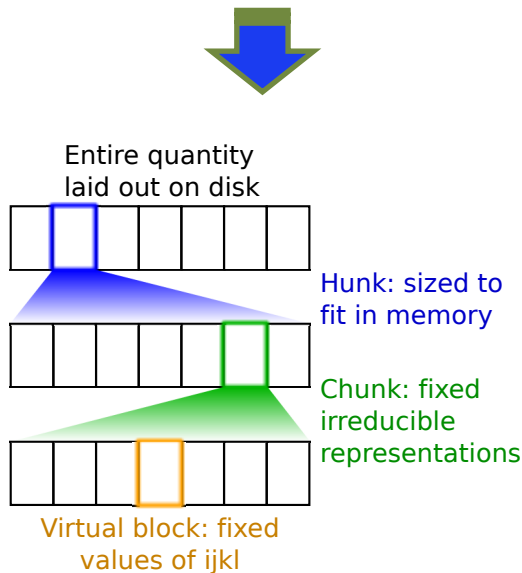


“Rank-k” MM and TC
on Xeon Phi 7210



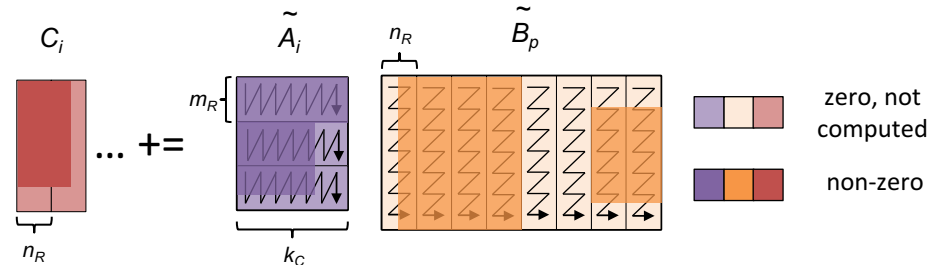
Quasi-Sparse Tensor Contractions

$$T_{i \leq j \leq k}^{abc}, R_{i \leq j \leq k \leq l}^{abcd}, W_{i \leq j \leq e}^{abc}, \text{ etc.}$$

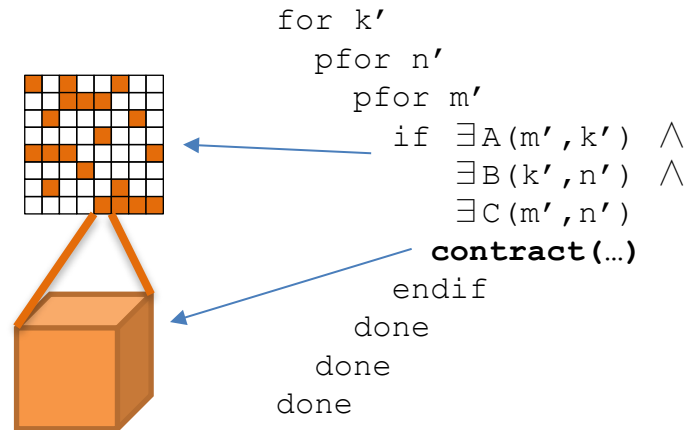


$$Z_{0,13,2,4}^{abcd} + = T_{0,5,21}^{abc} W_{13,2,4}^{5,21;d}$$

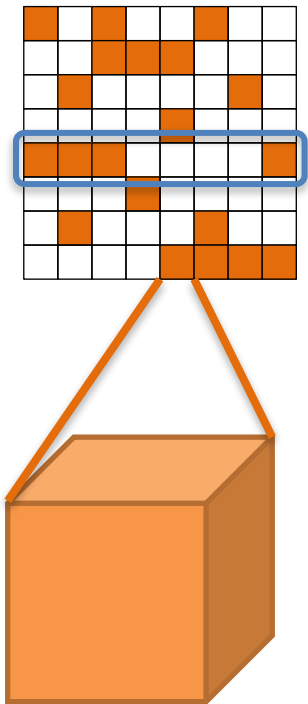
Option #1: Batch within TBLIS framework



Option #2: Batch outside of TBLIS framework

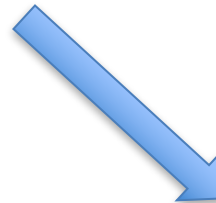


Quasi-Sparse Tensor Contractions



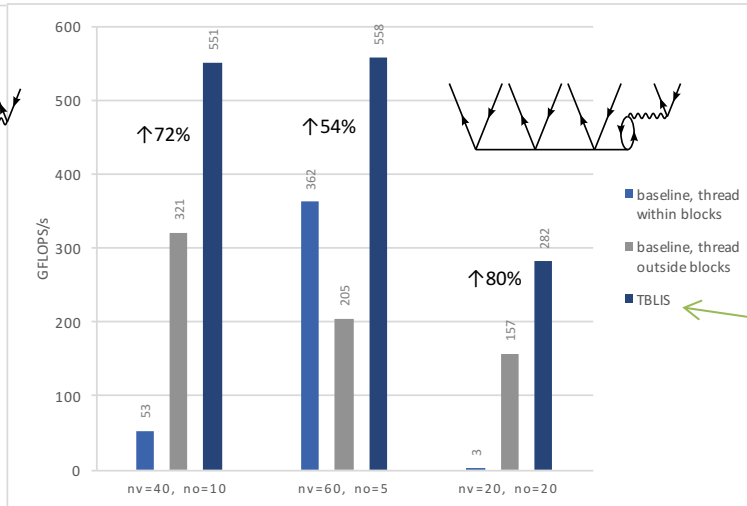
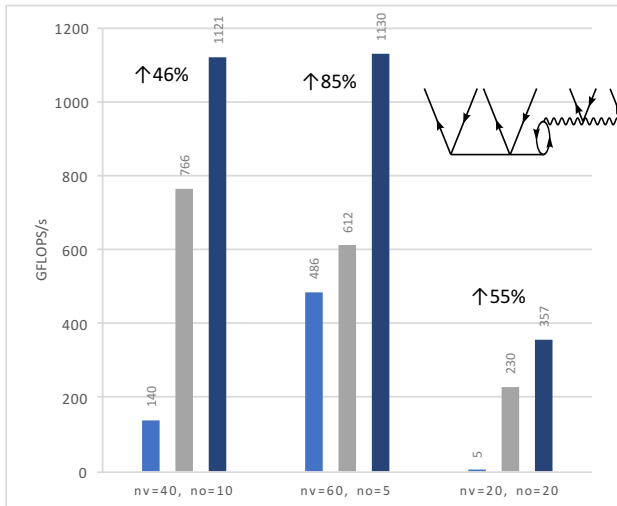
```
for k'  
  pfor n'  
    pfor m'  
      if  $\exists A(m', k') \wedge$   
         $\exists B(k', n') \wedge$   
         $\exists C(m', n')$   
        contract(...)  
      endif  
    done  
  done  
done
```

Use hierarchical
dynamic+static parallelism
and aggregate blocks when
possible.



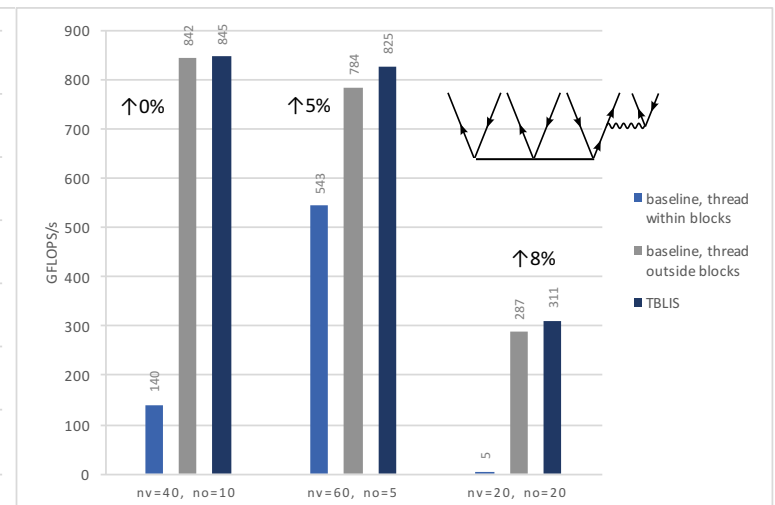
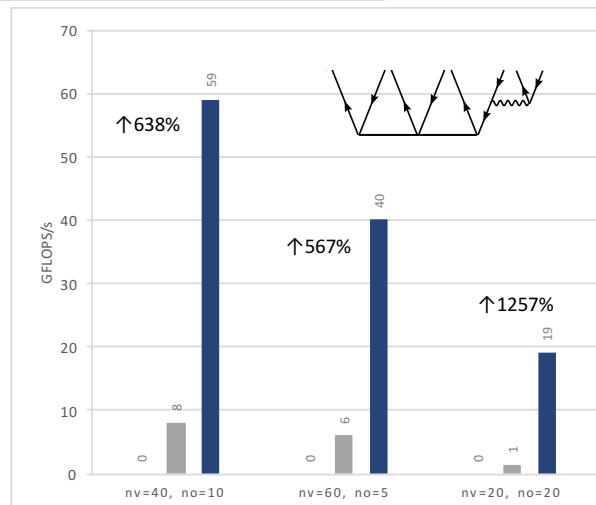
```
Split communicator into c_in & c_out  
pfor n' over c_out  
  pfor m' over c_out  
    ks = {}  
    for k'  
      if  $\exists A(m', k') \wedge$   
         $\exists B(k', n') \wedge$   
         $\exists C(m', n')$   
        append k' to ks  
      endif  
    done  
    pcontract(ks, ...) over c_in  
  done  
done
```

Quasi-Sparse Tensor Contractions

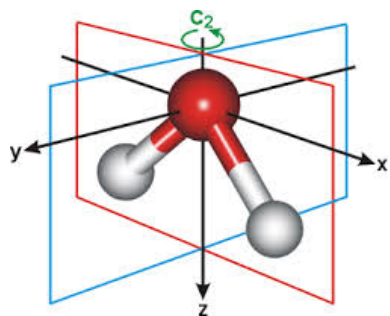


(uses TBLIS for inner tensor contraction)

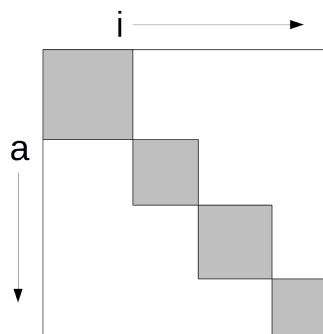
(adds hierarchical multithreading and block aggregation)



Taking Advantage of Structure



$$T_i^a \longrightarrow \Gamma_a \otimes \Gamma_i = \Gamma_T$$



Point Group Symmetry

Cost savings proportional to g^2 (g = number of irreducible representations/blocks).

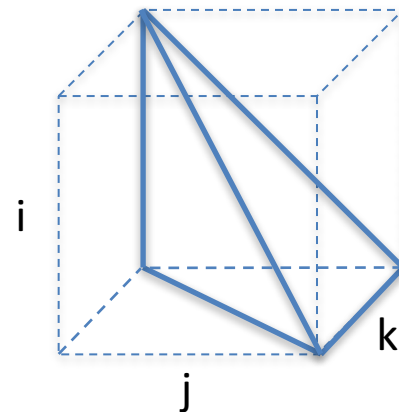
Permutational Symmetry

Factorial cost savings for increasing dimensionality.

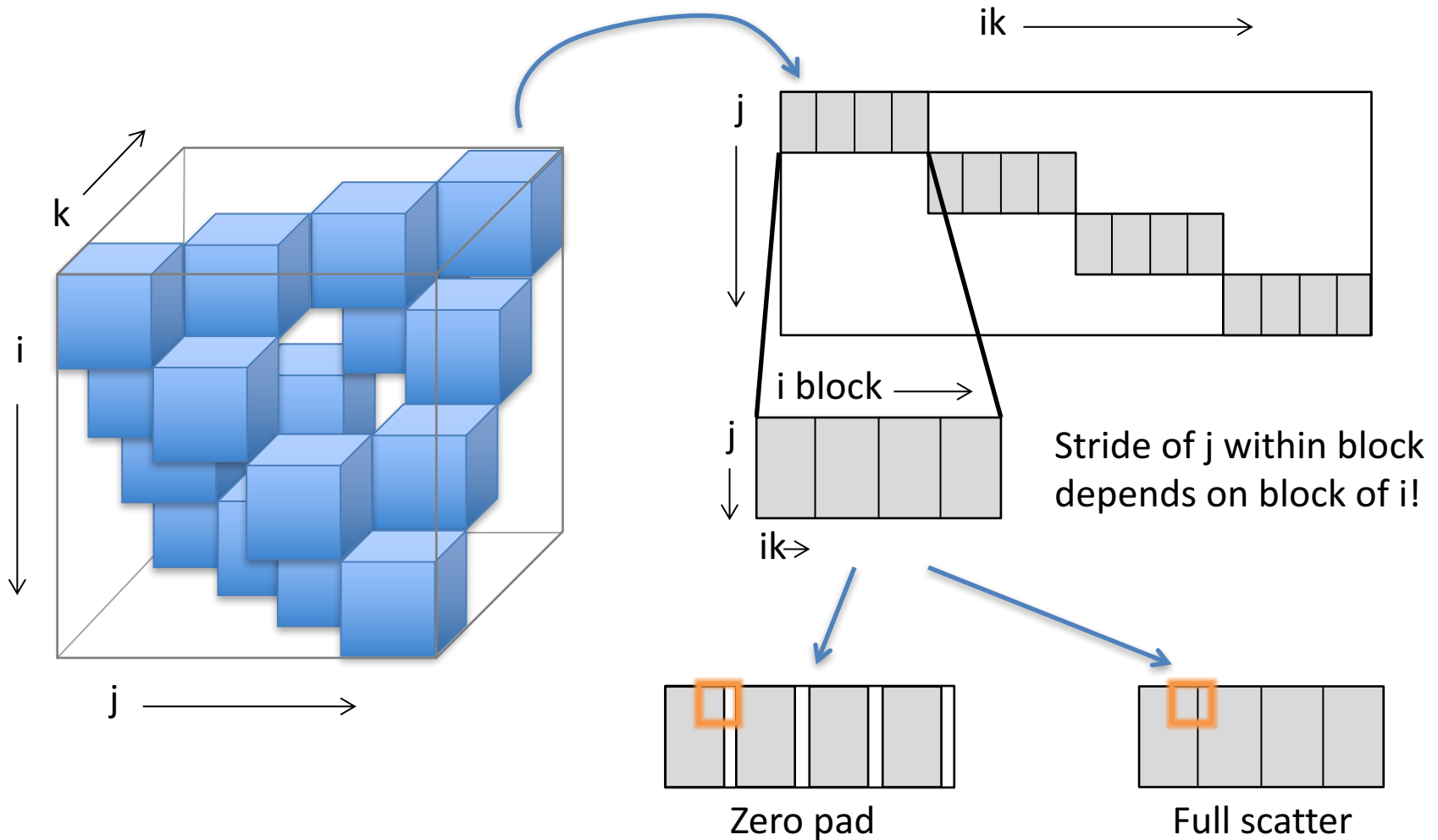
$$A_{ijk} = -A_{jik} = A_{jki} = -A_{kji} = A_{kij} = -A_{ikj}$$



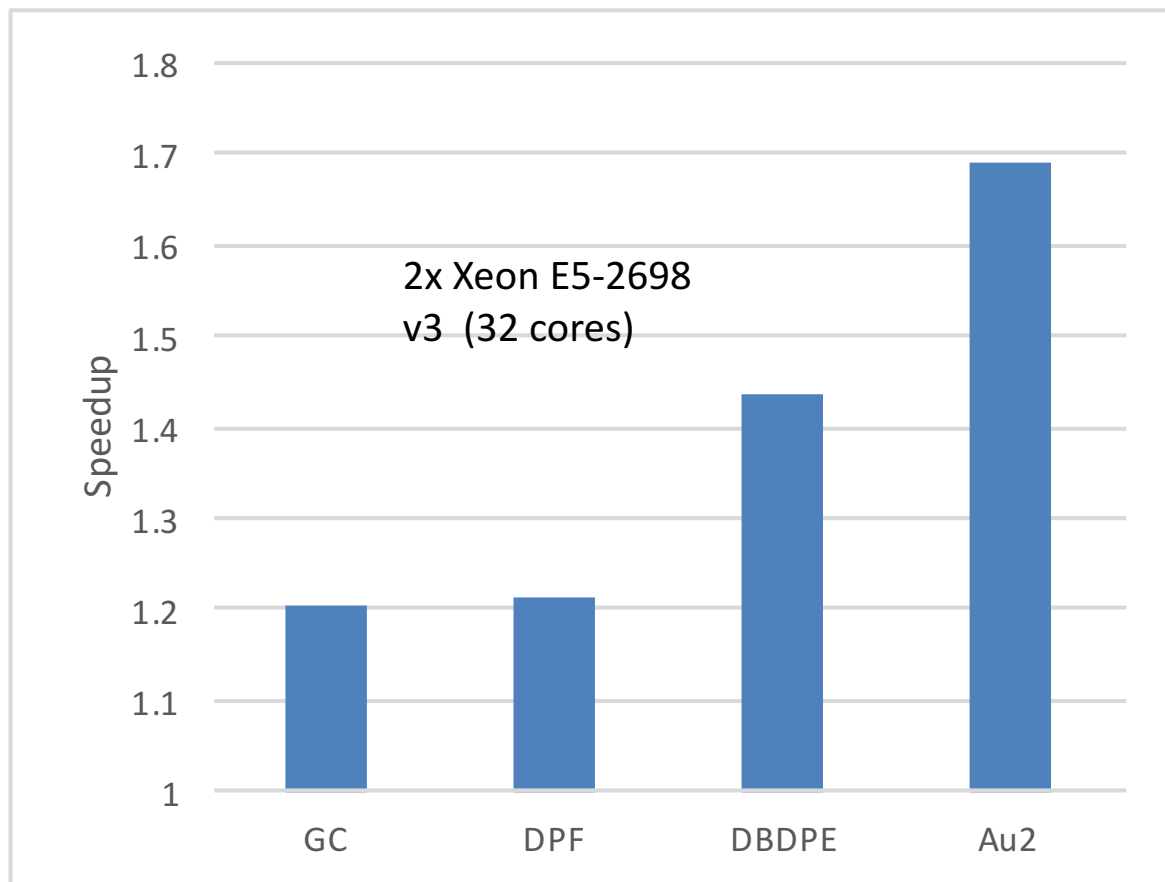
$$A_{i < j < k}$$



Taking Advantage of Structure



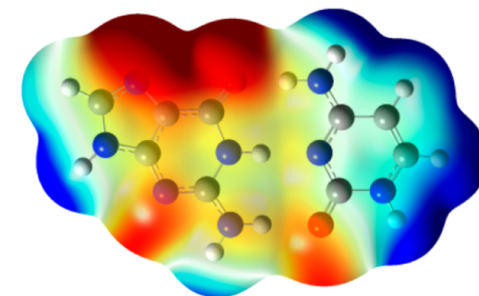
Speedup in computation of coupled cluster singles and doubles (CCSD) ground state energy when using TBLIS



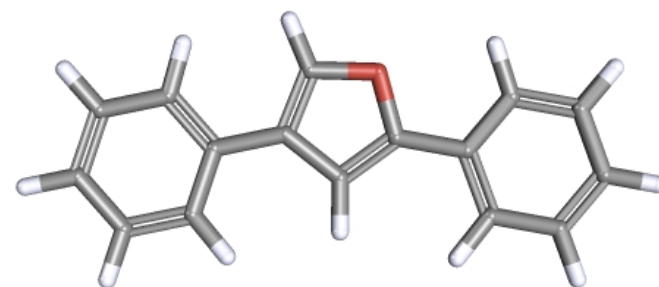
Less symmetry ← → More symmetry



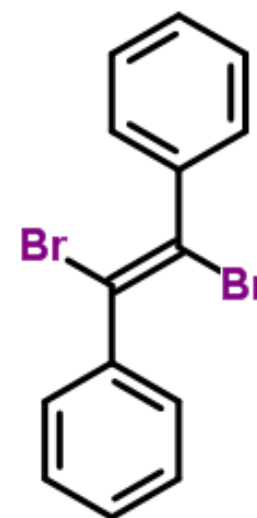
Gold dimer (**Au₂**)
all-electron, D_{2h} symmetry



Guanine-cytosine dimer (**GC**), no symmetry
Krepl et al., J. Phys. Chem. B 2013, 117, 1872



2,4-diphenylfuran (**DPF**), C_s symmetry

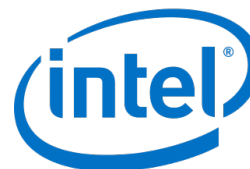


(E) 1,2-dibromo-1,2-diphenylethene (**DBDPE**)
planar, C_{2h} symmetry

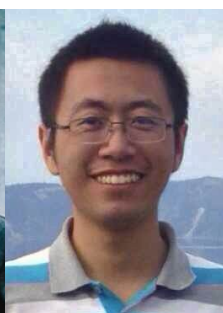
Summary

- Novel algorithms leveraging the **BLIS methodology** can significantly outperform DGEMM-based algorithms for **tensor contraction**.
- Breaking through the DGEMM barrier allows **new algorithms** to be implemented with high efficiency.

Thanks!



Robert van
de Geijn



Jianyu
Huang



Field
Van Zee



Tyler
Smith



Devangi
Parikh



#smallmoleculesmatter

