#### High Performance Computation and Interactive Display of Molecular Orbitals on GPUs and Multi-core CPUs

John Stone, Jan Saam, David Hardy, Kirby Vandivort, Wen-mei Hwu, Klaus Schulten

#### John Stone

#### Senior Research Programmer

Beckman Institute for Advanced Science and Technology University of Illinois at Urbana-Champaign http://www.ks.uiuc.edu/Research/gpu/



NIH Resource for Macromolecular Modeling and Bioinformatics

Second GPGPU Workshop, March 8, 2009

http://www.ks.uiuc.edu/

## VMD – "Visual Molecular Dynamics"

- Visualization and analysis of molecular dynamics simulations, sequence data, volumetric data, **quantum chemistry data**, particle systems
- User extensible with scripting and plugins
- http://www.ks.uiuc.edu/Research/vmd/





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# Molecular Orbitals

- Visualization of MOs aids in understanding the chemistry of molecular system
- MO spatial distribution is correlated with probability density for an electron



threonine



# Computing Molecular Orbitals

- Calculation of high resolution MO grids can require tens to hundreds of seconds in existing tools
- Existing tools cache MO grids as much as possible to avoid recomputation:
  - Doesn't eliminate the wait for initial calculation, hampers interactivity
  - Cached grids consume 100x-1000x more memory than MO coefficients







# Animating Molecular Orbitals

- Animation of (classical mechanics) molecular dynamics trajectories provides insight into simulation results
- To do the same for QM or QM/MM simulations one must compute MOs at ~10 FPS or more
- >100x speedup (GPU) over existing tools now makes this possible!





#### Molecular Orbital Computation and Display Process

One-time initialization

Read QM simulation log file, trajectory

Preprocess MO coefficient data eliminate duplicates, sort by type, etc...

For current frame and MO index,

retrieve MO wavefunction coefficients

For each trj frame, for each MO shown **Compute 3-D grid of MO wavefunction amplitudes** 

Most performance-demanding step, run on GPU...

Extract isosurface mesh from 3-D MO grid

Apply user coloring/texturing

and render the resulting surface



## CUDA Block/Grid Decomposition



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#### MO Kernel for One Grid Point (Naive C)

<pre>for (at=0; at<numatoms; at++)="" int="" prim_counter="atom_basis[at];&lt;/pre" {=""></numatoms;></pre>	Loop over atoms						
calc_distances_to_atom(&atompos[at], &xdist, &ydist, &zdist, &dist2, &xdiv);							
<pre>for (contracted_gto=0.0f, shell=0; shell &lt; num_shells_per_atom[at]; shell++) {     int shell_type = shell_symmetry[shell_counter];</pre>	Loop over shells						
<pre>for (prim=0; prim &lt; num_prim_per_shell[shell_counter]; prim++) {   float exponent = basis_array[prim_counter ];   float contract_coeff = basis_array[prim_counter + 1];   contracted_gto += contract_coeff * expf(-exponent*dist2);   prim_counter += 2; }</pre>	Loop over primitives: largest component of runtime, due to expf()						
<pre>for (tmpshell=0.0f, j=0, zdp=1.0f; j&lt;=shell_type; j++, zdp*=zdist) {     int imax = shell_type - j;     for (i=0, ydp=1.0f, xdp=pow(xdist, imax); i&lt;=imax; i++, ydp*=ydist, xdp*=xdiv)     tmpshell += wave_f[ifunc++] * xdp * ydp * zdp; }</pre>	Loop over angular momenta (unrolled in real code)						
<pre>value += tmpshell * contracted_gto; shell_counter++; }</pre>							
}							

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## Preprocessing of Atoms, Basis Set, and Wavefunction Coefficients

- Must make effective use of high bandwidth, lowlatency GPU on-chip memory, or CPU cache:
  - Overall storage requirement reduced by eliminating duplicate basis set coefficients
  - Sorting atoms by element type allows re-use of basis set coefficients for subsequent atoms of identical type
- Padding, alignment of arrays guarantees coalesced GPU global memory accesses, CPU SSE loads



### GPU Traversal of Atom Type, Basis Set, Shell Type, and Wavefunction Coefficients



- Loop iterations always access same or consecutive array elements for all threads in a thread block:
  - Yields good constant memory cache performance
  - Increases shared memory tile reuse



## Use of GPU On-chip Memory

- If total data less than 64 kB, use only const mem:
  - Broadcasts data to all threads, no global memory accesses!
- For large data, shared memory used as a programmanaged cache, coefficients loaded on-demand:
  - Tile data in shared mem is broadcast to 64 threads in a block
  - Nested loops traverse multiple coefficient arrays of varying length, complicates things significantly...
  - Key to performance is to locate tile loading checks outside of the two performance-critical inner loops
  - Tiles sized large enough to service entire inner loop runs
  - Only 27% slower than hardware caching provided by constant memory (GT200)



Array tile loaded in GPU shared memory. Tile size is a power-of-two, multiple of coalescing size, and allows simple indexing in inner loops (array indices are merely offset for reference within loaded tile).



Coefficient array in GPU global memory



#### VMD MO Performance Results for C<sub>60</sub> Sun Ultra 24: Intel Q6600, NVIDIA GTX 280

Kernel	Cores/GPUs	Runtime (s)	Speedup
CPU ICC-SSE	1	46.58	1.00
CPU ICC-SSE	4	11.74	3.97
CPU ICC-SSE-approx**	4	3.76	12.4
CUDA-tiled-shared	1	0.46	100.
CUDA-const-cache	1	0.37	126.
CUDA-const-cache-JIT*	1	0.27	173.
			(JIT 40% faster)

 $C_{60}$  basis set 6-31Gd. We used an unusually-high resolution MO grid for accurate timings. A more typical calculation has  $1/8^{th}$  the grid points.

\* Runtime-generated JIT kernel compiled using batch mode CUDA tools

\*\*Reduced-accuracy approximation of expf(), cannot be used for zero-valued MO isosurfaces





## VMD Orbital Dynamics Proof of Concept

One GPU can compute and animate this movie on-the-fly!

CUDA const-cache kernel, Sun Ultra 24, GeForce GTX 285

GPU MO grid calc.	<b>0.016 s</b>
CPU surface gen,	0.033 s
volume gradient,	
and GPU rendering	
<b>Total runtime</b>	<b>0.049</b> s
Frame rate	<b>20 FPS</b>



With GPU speedups over **100x**, previously insignificant CPU surface gen, gradient calc, and rendering are now **66%** of runtime. Need GPU-accelerated surface gen next...







#### Molecular Orbital Computation and Display Process Dynamic Kernel Generation, Just-In-Time (JIT) COmpilation

Read QM simulation log file, trajectory

Preprocess MO coefficient data

eliminate duplicates, sort by type, etc...

**Generate/compile basis set-specific CUDA kernel** 

For current frame and MO index,

retrieve MO wavefunction coefficients

For each trj frame, for each MO shown

**One-time** 

initialization

Compute 3-D grid of MO wavefunction amplitudes

using basis set-specific CUDA kernel

Extract isosurface mesh from 3-D MO grid

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Render the resulting surface

```
• • • • •
```

```
// loop over the shells belonging to this atom (or basis function)
for (shell=0; shell < maxshell; shell++) {
  float contracted_gto = 0.0f;</pre>
```

```
// Loop over the Gaussian primitives of this contracted
// basis function to build the atomic orbital
int maxprim = const_num_prim_per_shell[shell_counter];
int shell_type = const_shell_symmetry[shell_counter];
for (prim=0; prim < maxprim; prim++) {
  float exponent = const_basis_array[prim_counter ];
  float contract_coeff = const_basis_array[prim_counter + 1];
  contracted_gto += contract_coeff * exp2f(-exponent*dist2);
  prim_counter += 2;
```

```
}
```

```
/* multiply with the appropriate wavefunction coefficient */
float tmpshell=0;
switch (shell_type) {
   case S_SHELL:
    value += const_wave_f[ifunc++] * contracted_gto;
   break;
[.....]
```

#### case D\_SHELL:

```
tmpshell += const_wave_f[ifunc++] * xdist2;
tmpshell += const_wave_f[ifunc++] * ydist2;
tmpshell += const_wave_f[ifunc++] * zdist2;
tmpshell += const_wave_f[ifunc++] * xdist * ydist;
tmpshell += const_wave_f[ifunc++] * xdist * zdist;
tmpshell += const_wave_f[ifunc++] * ydist * zdist;
value += tmpshell * contracted_gto;
```

# Network for

<sup>b)</sup> General loop-based CUDA kernel

Dynamically-generated CUDA kernel (JIT) contracted\_gto = 1.832937 \* expf(-7.868272\*dist2); contracted\_gto += 1.405380 \* expf(-1.881289\*dist2); contracted\_gto += 0.701383 \* expf(-0.544249\*dist2); // P\_SHELL

tmpshell = const\_wave\_f[ifunc++] \* xdist; tmpshell += const\_wave\_f[ifunc++] \* ydist; tmpshell += const\_wave\_f[ifunc++] \* zdist; value += tmpshell \* contracted\_gto;

. . . . .

contracted\_gto = 0.187618 \* expf(-0.168714\*dist2); // S\_SHELL

 $value += const\_wave\_f[ifunc++] * contracted\_gto;$ 

contracted\_gto = 0.217969 \* expf(-0.168714\*dist2); // P\_SHELL

tmpshell = const\_wave\_f[ifunc++] \* xdist; tmpshell += const\_wave\_f[ifunc++] \* ydist; tmpshell += const\_wave\_f[ifunc++] \* zdist; value += tmpshell \* contracted\_gto;

contracted\_gto = 3.858403 \* expf(-0.800000\*dist2); // D\_SHELL tmpshell = const\_wave\_f[ifunc++] \* xdist2; tmpshell += const\_wave\_f[ifunc++] \* ydist2; tmpshell += const\_wave\_f[ifunc++] \* xdist \* ydist; tmpshell += const\_wave\_f[ifunc++] \* xdist \* ydist; tmpshell += const\_wave\_f[ifunc++] \* xdist \* zdist; tmpshell += const\_wave\_f[ifunc++] \* ydist \* zdist; tmpshell += const\_wave\_f[ifunc++] \* ydist \* zdist;

#### Performance Evaluation: Molekel, MacMolPlt, and VMD Sun Ultra 24: Intel Q6600, NVIDIA GTX 280

			C <sub>60</sub> -A	C <sub>60</sub> -B	Thr-A	Thr-B	Kr-A	Kr-B
	Atoms		60	60	17	17	1	1
	Basis funcs (unique)		300 (5)	900 (15)	49 (16)	170 (59)	19 (19)	84 (84)
Kernel		Cores GPUs	Speedup vs. Molekel on 1 CPU core					
Molekel		1*	1.0 1.0 1.0 1.0 1.0 1.				1.0	
MacMo	lPlt	4	2.4 2.6 2.1 2.4 4.3		4.5			
VMD G	CC-cephes	4	3.2 4.0 3.0 3.5 4.3		6.5			
VMD IO	CC-SSE-cephes	4	16.8 17.2 13.9 12.6 17.3 2		21.5			
VMD IO	CC-SSE-approx**	4	59.3	53.4	50.4	49.2	54.8	69.8
VMD C	UDA-const-cache	1	552.3	533.5	355.9	421.3	193.1	571.6



## Future Work

- Tune Multi-GPU implementation to workaround small kernel launch delays that adversely impact animation speed
- Further development of runtime-generated MO kernels using new CUDA JIT compilation APIs
- Multi-pass approach with spatial decomposition and distance-based cutoff to truncate rapidly decaying exponentials (CPU+GPU cooperation)



## Acknowledgements

- Theoretical and Computational Biophysics Group, IMPACT group, University of Illinois at Urbana-Champaign
- CUDA team at NVIDIA
- NIH support: P41-RR05969

