## SCF optimizations

## Execution times of 16 atom test

| Optimization | Time $(\mathbf{s})$ | Speedup |
| :--- | :--- | :--- |
| Reference implementation | 176.35 | 1 |
| Exploit symmetry of g() | 30.58 | 5.8 |
| Use lookup tables in g() | 19.06 | 9.3 |
| BLAS and LAPACK matrix routines | 18.06 | 9.8 |
| Don't recompute h() each iteration | 17.95 | 9.8 |
| Cache results of g() | 9.45 | 18.7 |

- All tests were performed on a workstation with an Intel Xeon processor running at 2.67 GHz .
- Each optimization test includes all optimizations listed above it.
- Caching $g()$ leads to the best performance, but the size of the cache grows with $\mathrm{N}^{4}$.


## Additional work skipping test

```
double twoel(double schwmax) {
    int i, j, k, l;
    for (i = 0; i < nbfn; i++) {
    for (j = 0; j < nbfn; j++) {
        if ((g_schwarz[i][j] * schwmax) < tol2e)
            {icut1 += nbfn * nbfn; continue;}
        double KLTest = tol2e / g_schwarz[i][j];
    for (k = 0; k < nbfn; k++) {
    for (l = 0; l < nbfn; l++) {
        if (g_schwarz[k][l] < KLTest)
            {icut2 ++; continue;}
        icut3 ++;
        double gg = g(i, j, k, l);
        g_fock[i][j] += ( gg *
        g_dens[k][l]);
        g_fock[i][k] -= (0.50 * gg *
        g_dens[j][l]);
    } } } }
    return (0.50 * contract_matrices(g_fock,
        g_dens));
}
        --
```

- The inner-most loop always tests if g_schwarz [k] [l] is less than KLTest
- While creating g_schwarz, we can also determine what the maximum g_schwarz [ $k$ ] [l] is for each column $k$ and store it in

```
g_schwarz_max_row_value[k]
```

- Then, in the $k$ loop, we can check if g_schwarz_max_row_value [k] is less than KLTest
- If it is, then we know that all the work in the I loop will be skipped, and can skip that loop entirely
- The additional memory requirements for this are small as we only need a onedimensional array of size nbfn


## Code changes required for additional work skipping test

```
double twoel(double schwmax) {
    int i, j, k, l;
    for (i = 0; i < nbfn; i++) {
    for (j = 0; j < nbfn; j++) {
        if ((g_schwarz[i][j] * schwmax) < tol2e)
            {icut1 += nbfn * nbfn; continue;}
        double KLTest = tol2e / g_schwarz[i][j];
    for (k = 0; k < nbfn; k++) {
        if (g_schwarx_max_row_value[k] < KLTest)
            {icut4 ++; continue;}
        for (l = 0; l < nbfn; l++) {
        if (g_schwarz[k][l] < KLTest)
            {icut2 ++; continue;}
        icut3 ++;
        double gg = g(i, j, k, l);
        g_fock[i][j] += ( gg *
        g_dens[k][l]);
        g_fock[i][k] -= (0.50 * gg *
        g_dens[j][l]);
    } } } }
    return (0.50 * contract_matrices(g_fock,
        g_dens));
}
```

```
double makesz() {
    int i, j;
    double smax = 0.0;
    for (i = 0; i < nbfn; i++) {
double row_max = 0.0; 
        double gg = sqrt( g(i, j, i, j) );
        if (gg > smax) smax = gg;
        g_schwarz[i][j] = gg;
        if (gg > row_max) row_max = gg;
    }
    g_schwarx_max_row_value[i] = row_max;
    }
    return smax;
}

\section*{Exploit symmetry in g()}
- \(g(i, j, j, \mathrm{l})\) returns the same value in the following cases:
- i and j are swapped
- k and I are swapped
- i,j and k,l are swapped
- We can use this to reduce the calls to \(g()\) to \(1 / 8\) th of the original.

\section*{Exploit symmetry in g()}
- update(a, b, c, d, gg) \{
```

g_fock[b][a] += (gg * g_dens[c][d]);
g_fock[c][a] -= (0.50 * gg * g_dens[b][d]);}

```
- update() should be an inline function or a macro to minimize overhead.
- We can do \(g g=g(i, j, k, l)\) once then perform the following updates:
- update(i, j, k, l, gg)
- update(i, j, I, k, gg)
- update(j, i, k, l, gg)
- update(j, i, I, k, gg)
- update(k, l, i, j, gg)
- update(l, k, i, j, gg)
- update(k, l, j, i, gg)
- update(l, k, j, i, gg)

\section*{Exploit symmetry in g()}
- There are special cases to consider:
- We must be sure to only iterate over i,j,k,I values which we haven't already processed due to the symmetry.
- If \(\mathrm{i}=\mathrm{j}\), we can't perform updates which swap i and j . Doing so would lead to the wrong values being accumulated into various g_fock entries.
- In addition to \(\mathrm{i}=\mathrm{j}\), we must also take care when \(\mathrm{k}=\mathrm{l}\) or \(i, j=k, l\) as well as any combinations of them.

\section*{Exploit symmetry in g()}
- We end up with 6 separate variants of the twoel loop:
- i, j, k, I can be freely swapped
- \(\mathrm{i}=\mathrm{j}\)
- \(k=1\)
- \(\mathrm{i}, \mathrm{j}=\mathrm{k}, \mathrm{l}\)
- \(\mathrm{i}=\mathrm{j}\) and \(\mathrm{k}=\mathrm{I}\)
- \(\mathrm{i}=\mathrm{j}=\mathrm{k}=\mathrm{l}\)

\section*{Exploit symmetry in g()}
- Loop structure when i, j, k, I can be freely swapped:
```

for (i = 0; i < nbfn; i++) {
for (j = i + 1; j < nbfn; j++) {
for (k = i; k < nbfn; k++) {
if (k == i) l_start = 1 + j;
else l_start = 1 + k;
for (l = l_start; l < n.bfn; l++) {
//calculate g(i,j,k,l);
//perform all eight updates
}}}}

```
- The starting index for the 1 loop is a special case which depends on whether \(k\) is equal to \(i\)
- The other loops can be easily created by removing the loop and index for the appropriate symmetry
- The number of updates to perform will be fewer since one or more of the swaps will not alter the arguments
- The special case for the initial value of \(l\) is only required in the fully symmetric case. In all other cases, the 1 loop is either removed, or the initial value reduces to \(1+k\)
- All other variants have 1 set equal to another index:
- \(k=1\)
- \(\mathrm{i}, \mathrm{j}=\mathrm{k}, \mathrm{l}\)
- \(\mathrm{i}=\mathrm{j}\) and \(\mathrm{k}=1\)
- \(i=j=k=1\)
- All loops could potentially be combined into one, but doing so may hurt performance.
- Conditional updates in the inner loop would create a lot of overhead.

Other methods might cause difficulties when trying to parallelize the code or could make automatic compiler optimizations impossible.

\section*{Precomputing lookup table for g()}
- Many calculations in g() only rely on i,j or k,l.
- Intermediate values are calculated for both ij and kl inputs, then later used together.
- All calculations which only rely on one of the pairs can easily be computed during initialization and stored in an \(\mathrm{N}^{2}\) sized array.

\section*{Removing \(\exp ()\) function from \(g()\)}

\section*{Calculation of exijkI}
```

double dxij = x[i] - x[j];
double dyij = y[i] - y[j];
double dzij = z[i] - z[j];
double dxkl = x[k] - x[l];
double dykl = y[k] - y[l];
double dzkl = z[k] - z[l]
double rab2 = dxij * dxij + dyij * dyij + dzij * dzij;
double rcd2 = dxkl * dxkl + dykl * dykl + dzkl * dzkl;
double expntIJ = expnt[i] + expnt[j];
double expntKL = expnt[k] + expnt[l];
double facij = expnt[i] * expnt[j] / expntIJ;
double fackl = expnt[k] * expnt[l] / expntKL;
double exijkl = exprjh(-facij * rab2 - fackl * rcd2);

```

\section*{Simplifying exijkl calculation}
```

$r_{i j}^{2}=\left(x_{i}-x_{j}\right)^{2}+\left(y_{i}-y_{j}\right)^{2}+\left(z_{i}-z_{j}\right)^{2}$
expnt $_{i j}=$ expnt $_{i}+$ expnt $_{j}$
fac $_{i j}=\frac{\text { expnt }_{i} * \text { expnt }_{j}}{\text { expnt }_{i j}}$
$e x_{i j k l}=e^{\left(-f a c_{i j} * r_{i j}{ }^{2}\right)+\left(-f a c_{k l} * r_{k l}{ }^{2}\right)}$
$e x_{i j k l}$ can be substituted with:
$e x_{i j k l}=e^{\left(-f a c_{i j} * r_{i j}^{2}\right)} * e^{\left(-f a c_{k l} * r_{k l}^{2}\right)}$

```

Then, we can define \(e x_{i j}\) as:
\(e x_{i j}=e^{\left(-f a c_{i j} * r_{i j}{ }^{2}\right)}\)
Now we can express ex \(x_{i j k l}\) as the product of two terms, each of which only depends on ij or kl.
\(e x_{i j k l}=e x_{i j} * e x_{k l}\)
So by storing a table of all possible values for \(e x_{i j}\), we can replaced 16 memory lookups, 10 multiplications, 13 additions/subtractions, 2 divisions and an \(e^{x}\) calculation with two \({ }^{1}\) memory lookups and a single multiplication.

\footnotetext{
\({ }^{1}\) There will be two additional memory lookups required to get expnt \({ }_{i j}\) and expnt \(_{k l}\) for other calculations in g 0
}

\section*{Precomputing lookup table for \(\mathrm{g}(\) )}
- Six values can be precomputed:
- ex \(x_{i j}\)
- expnt \({ }_{i j}\)
- \(x_{i j}\) (Replaces xp and xq )
- \(y_{i j}\) (Replaces yp and yq)
- \(z_{i j}\) (Replaces zp and zq)
- rnorm \({ }_{i j}\)
- The same lookup tables are use for both i,j and k,l indicies
- Requires additional \(6 * N^{2}\) of memory
- Cache performance can be improved by storing all precomputed values as an array of structs rather than in separate arrays.

\section*{Execution time breakdown as problem size increases}

- As the problem size increases, the time spent diagonalizing the matrix (Eigen_std and dgemm in above chart) becomes a larger portion of the overall execution time
- Fortunately, Eigen_std and dgemm can be replaced with standard matrix operations from BLAS and LAPACK

\section*{Replacement Matrix Routines}
- BLAS can provide replacements for the following operations:
- dgemm() calls in diagon()
- ddot() can replace contract_matricies()
- dscal() and daxpy() can replace damp()
- cblas_dscal(nbfn*nbfn, fac, g_dens, 1);
- // g_dens = fac * g_dens
- cblas_daxpy(nbfn*nbfn, 1.0 - fac, g_work, 1, g_dens, 1);
- The resulting calculation is:
- Which is identical to the original operation in damp()
- This may not increase performance if the compiler was already able to vectorize damp() since using BLAS routines requires two iterations over the array instead of one.
- LAPACK routines can replace the functions in diagonalize.c:
- rsg() \(\rightarrow\) dsygv()
- \(\mathrm{rs}(\) ) \(\rightarrow\) dsyev()

\section*{Removing oneel()}
- The oneel() function doesn't take long to run, but still performs unnecessary calculations.
- g_fock is initialized with the same values each iteration.
- Instead, we can save a copy of the initial g_fock values and copy it back each iteration.
- We still have to call contract_matricies() to determine the one-electron energy contribution.

\section*{Execution time comparison}

Unoptimized dgemm, Eigen_std and oneel


Optimized oneel with dgemm and Eigen_std using BLAS and LAPACK

161820222426283032343638404244464850 Number of atoms

■ Eigen_std dgemm ■ twoel onel

\section*{Conventional method vs. direct method}

- Both conventional and direct methods included all optimizations.
- The conventional method is faster than the direct method in all tests performed.
- However, the usefulness of the conventional method is limited due to the amount of memory required, the additional power requirements, and the decreasing performance as the problem size increases.

\section*{Summary}
- \(g()\) has an 8 -way symmetry which can be exploited to reduce required calculations.
- Many intermediate calculations in g() can be precomputed and stored in a lookup table.
- Allows removing a slow \(\mathrm{e}^{\mathrm{x}}\) calculation.
- Reduces number of memory operations required in g() .
- Matrix and vector routines can be trivially replaced with optimized libraries.
- Conventional method can speedup \(g()\) even further.
- Very high memory requirements.
- Incremental speedup with other optimizations only \(\sim 2 x\).```

