

The basics of the DMC code

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What can it do?

Quantities of interest

- Ground state energies and densities.
- Energy distributions.

Implemented Systems

- Harmonic oscillator systems (2D, 3D, doublewells)
- Atomic systems (Atoms, homonuclear diatomic molecules)

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Underlying goals and assumptions

In a nutshell

Ab-initio, Efficiency, Transparency

Specifics: Assumptions

- Single Slater-determinant ansatz.
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Compromise to ensure efficiency and “ideal scaling”

SP-basis \rightarrow Det. Basis \nrightarrow ~~Combine Determinants~~

Start from the bottom up

- Implement single particle wave functions $\Phi_j(\vec{r}_i)$.

```
class BasisFunctions {  
public:  
    BasisFunctions();  
  
    virtual double eval(const Walker* walker, int i) = 0;  
};
```

```
//n_x = n_y = n_z = 1
double HarmonicOscillator3D_15::eval(const Walker* walker, int i) {

    x = walker->r(i, 0);
    y = walker->r(i, 1);
    z = walker->r(i, 2);

    //x*y*z*exp(-k^2*r^2/2)

    H = x*y*z;
    return H*(exp_factor);

}
```

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```
class Orbitals {
protected:

    BasisFunctions** basis_functions;
    BasisFunctions*** del_basis_functions;
    BasisFunctions** lapl_basis_functions;

    //Important: All basis elements and the orbital wrapper
    //SHARE parameter references
    virtual void set_parameter(double parameter, int n) = 0;

public:

    Orbitals(int n_p, int dim);

    virtual double      phi(const Walker* walker, int i, int q);
    virtual double      del_phi(const Walker* walker, int i, int q, int d);
    virtual double      lapl_phi(const Walker* walker, int i, int q);

};
```

```
double Orbitals::phi(const Walker* walker, int particle, int q_num) {  
    return basis_functions[q_num]->eval(walker, particle);  
}  
  
double Orbitals::del_phi(const Walker* walker, int particle, int q_num, int d)  
    return del_basis_functions[d][q_num]->eval(walker, particle);  
}  
  
double Orbitals::lapl_phi(const Walker* walker, int particle, int q_num) {  
    return lapl_basis_functions[q_num]->eval(walker, particle);  
}
```

Idea: Fill the basis function vectors in the subclass constructor,
and you're good to go.

```
HarmonicOscillator::HarmonicOscillator{  
    ...  
  
    basis_functions[0] = new HarmonicOscillator3D_0(...);  
    basis_functions[1] = new HarmonicOscillator3D_1(...);  
    basis_functions[2] = new HarmonicOscillator3D_2(...);  
    ...  
  
    del_basis_functions[0][0] = new del_HarmonicOscillator3D_0_x(...);  
    del_basis_functions[1][0] = new del_HarmonicOscillator3D_0_y(...);  
    del_basis_functions[2][0] = new del_HarmonicOscillator3D_0_z(...);  
    del_basis_functions[0][1] = new del_HarmonicOscillator3D_1_x(...);  
    del_basis_functions[1][1] = new del_HarmonicOscillator3D_1_y(...);  
    del_basis_functions[2][1] = new del_HarmonicOscillator3D_1_z(...);  
    ...  
  
    lapl_basis_functions[0] = new lapl_HarmonicOscillator3D_0(...);  
    lapl_basis_functions[1] = new lapl_HarmonicOscillator3D_1(...);  
    ...  
}
```

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Expanded Bases

```
class ExpandedBasis : public Orbitals {
public:
    ExpandedBasis(...);

    double phi(...);
    double del_phi(...);
    double lapl_phi(...);

protected:
    arma::mat coeffs;

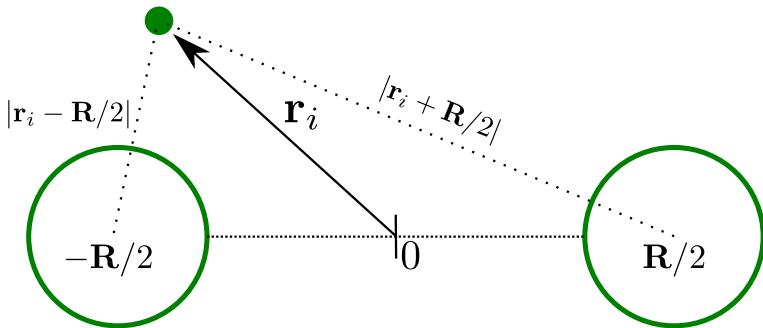
    Orbitals* basis;

};
```

Expanded Bases

```
double ExpandedBasis::phi(const Walker* walker, int i, int q) {  
  
    double value = 0;  
  
    //Dividing basis_size by half assuming a two-level system.  
    for (int m = 0; m < basis_size / 2; m++) {  
        value += coeffs(q, m) * basis->phi(walker, i, m);  
    }  
  
    return value;  
  
}
```

Can be loaded into the QMC machinery as any other Orbital instance.



$$\hat{\mathbf{H}}_{\text{Mol.}}(\mathbf{r}, \mathbf{R}) = \sum_{i=1}^N \left[-\frac{1}{2} \nabla_i^2 - \frac{Z}{|\mathbf{r}_i + \mathbf{R}/2|} - \frac{Z}{|\mathbf{r}_i - \mathbf{R}/2|} \right] + \frac{Z^2}{R} + \sum_{i < j} \frac{1}{r_{ij}}.$$

(di)Molecules

```
double DiAtomCore::get_pot_E(const Walker* walker) const {  
  
    double e_pot = 0;  
    double com_corr, shared;  
  
    double quarterR2 = 0.25*(*R)*(*R);  
    for (int i = 0; i < n_p; i++) {  
  
        shared = walker->get_r_i2(i)+ quarterR2;  
        com_corr = (*R)*walker->r(i, 0);  
  
        e_pot -= Z*(1./sqrt(shared + com_corr) + 1./sqrt(shared - com_corr));  
    }  
  
    e_pot += Z*Z/(*R);  
  
    return e_pot;  
}
```

(di)Molecules

```
class DiTransform : public Orbitals {  
...  
  
protected:  
    double* R;  
  
    Orbitals* nucleus1, nucleus2;  
    Walker* walker_nucleus1, walker_nucleus2;  
  
    //Wrap wrap wrap, it's christmas!  
    double get_parameter(int n) {  
        return nucleus1->get_parameter(n);  
    }  
  
    void set_parameter(double parameter, int n) {  
        nucleus1->set_parameter(parameter, n);  
        nucleus2->set_parameter(parameter, n);  
    }  
};
```

(di)Molecules

- A Walker instance for each nucleus ensures minimal overhead when transforming from atoms to molecules (precalculates distances etc.).
- An Orbital instance for each nucleus ensures that all optimizations from the single atom scheme can be carried to the molecular one (precalculates exponential factors etc.).

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(di)Molecules

```
void DiTransform::set_qnum_indie_terms(Walker* walker, int i) {  
  
    walker->calc_r_i(i);  
  
    //Apply the molecular transformation!  
    walker_nucleus1->r.row(i) = walker->r.row(i);  
    walker_nucleus2->r.row(i) = walker->r.row(i);  
  
    walker_nucleus1->r(i, 0) += (*R) / 2;  
    walker_nucleus2->r(i, 0) -= (*R) / 2;  
  
    double shared = walker->get_r_i2(i) + 0.25 * (*R)*(*R);  
    double comm_spec = walker->r(i, 0)*(*R);  
  
    walker_nucleus1->r2(i) = shared + comm_spec;  
    walker_nucleus2->r2(i) = shared - comm_spec;  
  
    nucleus1->set_qnum_indie_terms(walker_nucleus1, i);  
    nucleus2->set_qnum_indie_terms(walker_nucleus2, i);  
}
```


Transforming SPWFs to molecular SPWFs

Plus and minus refers to the two nuclei, H refers to the standard hydrogen-like basis.

$$\begin{aligned}\phi_{nlm}^+(\mathbf{r}_i, \mathbf{R}) &= \phi_{nlm}^H(\mathbf{r}_i + \mathbf{R}/2) + \phi_{nlm}^H(\mathbf{r}_i - \mathbf{R}/2), \\ \phi_{nlm}^-(\mathbf{r}_i, \mathbf{R}) &= \phi_{nlm}^H(\mathbf{r}_i + \mathbf{R}/2) - \phi_{nlm}^H(\mathbf{r}_i - \mathbf{R}/2),\end{aligned}$$

which reads “electron surrounding first nucleus combined with electron surrounding second nucleus”.

Applying the transformation

```
double DiTransform::phi(const Walker* walker, int i, int q) {  
  
    (void) walker;  
    int sign = minusPower(q);  
  
    return nucleus1->phi(walker_nucleus1, i, q / 2) +  
           sign * nucleus2->phi(walker_nucleus2, i, q / 2);  
  
}
```

We can reuse closed form expressions

$$\begin{aligned}
 \mathbf{j} \cdot \nabla_i \phi_{nlm}^{\pm}(\mathbf{r}_i, \mathbf{R}) &= \underbrace{\frac{\partial(y_i + R_y/2)}{\partial y_i}}_1 \frac{\partial \phi_{nlm}^H(\mathbf{r}_i + \mathbf{R}/2)}{\partial(y_i + R_y/2)} \\
 &\pm \underbrace{\frac{\partial(y_i - R_y/2)}{\partial y_i}}_1 \frac{\partial \phi_{nlm}^H(\mathbf{r}_i - \mathbf{R}/2)}{\partial(y_i - R_y/2)} \\
 &= \frac{\partial \phi_{nlm}^H(\mathbf{r}_i + \mathbf{R}/2)}{\partial(y_i + R_y/2)} \pm \frac{\partial \phi_{nlm}^H(\mathbf{r}_i - \mathbf{R}/2)}{\partial(y_i - R_y/2)} \\
 &= \frac{\partial \phi_{nlm}^H(\tilde{\mathbf{R}}_i^+)}{\partial \tilde{Y}_i^+} \pm \frac{\partial \phi_{nlm}^H(\tilde{\mathbf{R}}_i^-)}{\partial \tilde{Y}_i^-},
 \end{aligned}$$

We can reuse closed form expressions

```
double DiTransform::lapl_phi(const Walker* walker, int particle, int q_num) {  
  
    (void) walker;  
    int sign = minusPower(q_num);  
  
    return nucleus1->lapl_phi(walker_nucleus1, particle, q_num / 2) +  
           sign * nucleus2->lapl_phi(walker_nucleus2, particle, q_num / 2);  
}
```

Generalizable to N-atomic molecules?

YES.

Challenges:

- Current model breaks down around O_2 .
- Need a better trial wave function.
- Multi-determinants are out of the question.
- More advanced Jastrow is out of the question.
- Expanded single particle states are gold.
- Slater orbitals are gold.

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Molecule	R	E_{VMC}	E_{DMC}	Expt.
H ₂	1.4	-1.1551(3)	-1.1745(3)	-1.1746
⋮				
O ₂	2.282	-143.97(2)	-148.53(2)	-150.3268

Table: Refs. for R and Expt.: [3, 2, 1].

Parametrizations

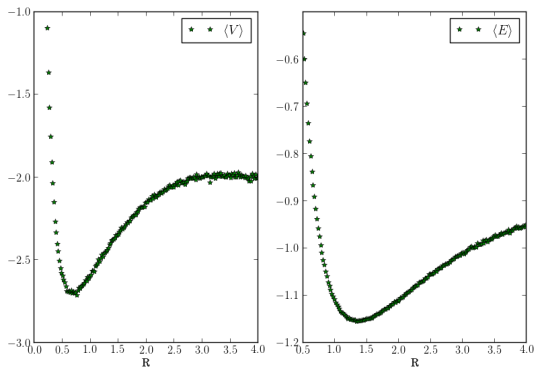


Figure: H_2

Parametrizations

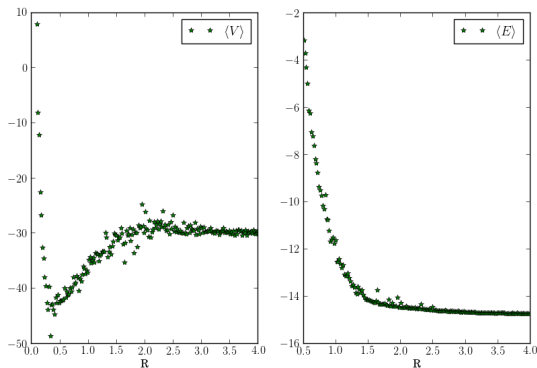


Figure: Li_2



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