# Recent progresses in the variational reduced-density-matrix method

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- Introduction of the RDM method.
- Recent results.
- Some open problems.



#### Introduction of the RDM method.

#### The RDM method: 2-RDM as basic variable

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Equivalent to the Schrödinger equation Ground state energy: Minimize directly!

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Equivalent to the Schrödinger equation Ground state energy: Minimize directly! *N*-representability condition; the only one approximation Our goal: doing chemistry from the first principle, faster calculation and deeper understanding

# Our target

- ab initio...theoretically and practically good approximation
- faster method ...mathematically simpler
- deeper understanding...electronic structure

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$$H = \sum_{ij} v_j^i a_i^{\dagger} a_j + \frac{1}{2} \sum_{i_1 i_2 j_1 j_2} w_{j_1 j_2}^{i_1 i_2} a_{i_1}^{\dagger} a_{i_2}^{\dagger} a_{j_2} a_{j_1}$$

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The ground state energy becomes...

$$E_g = \min \langle \Psi | H | \Psi \rangle$$
  
= 
$$\min \sum_{ij} v_j^i \langle \Psi | a_i^{\dagger} a_j | \Psi \rangle + \frac{1}{2} \sum_{i_1 i_2 j_1 j_2} w_{j_1 j_2}^{i_1 i_2} \langle \Psi | a_{i_1}^{\dagger} a_{i_2}^{\dagger} a_{j_2} a_{j_1} | \Psi \rangle$$
  
= 
$$\min \{ \sum_{ij} v_j^i \gamma_j^i + \sum_{i_1 i_2 j_1 j_2} w_{j_1 j_2}^{i_1 i_2} \Gamma_{j_1 j_2}^{i_1 i_2} \}$$

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Definition of 1, 2-RDMs

$$\Gamma^{i_1i_2}_{j_1j_2}=\frac{1}{2}\langle\Psi|a^{\dagger}_{i_1}a^{\dagger}_{i_2}a_{j_2}a_{j_1}|\Psi\rangle, \ \gamma^i_j=\langle\Psi|a^{\dagger}_ia_j|\Psi\rangle.$$

#### N-representability condition

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$$E_g = \min_{\mathcal{P}} \{ \sum_{ij} v_j^i \gamma_j^i + \sum_{i_1 i_2 j_1 j_2} w_{j_1 j_2}^{i_1 i_2} \Gamma_{j_1 j_2}^{i_1 i_2} \}$$

 $\gamma, \Gamma \in \mathcal{P}$  should satisfy *N*-representability condition:

$$\Gamma(12|1'2') \to \Psi(123\cdots N)$$

$$\gamma(1|1') \rightarrow \Psi(123 \cdots N).$$

Encodes two-body effects completely. Very compact.

Approximation (necessary) condition : where Physics and Chemistry are

• P, Q-condition, ensemble 1-RDM condition [Coleman 1963]

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- Construction of 2-particle density [Pistol 2004, 2006]



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#### Can evaluate total energy exactly via 1 and 2-RDM

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- Can evaluate total energy exactly via 1 and 2-RDM
- only one approximation is N-representability condition (aka theory of everything)

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Method	# of variable (discritized)	Exact?
Ψ	N, ( <b>r</b> !)	Yes
Γ(12 1'2')	$4, (r^4)$	Yes

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Γ(12 1'2')	<b>4</b> , ( <b>r</b> <sup>4</sup> )	Yes

Do not depend on the size of the system Equivalent to Schrödinger eq. (ground state)

# Mathematically simpler: minimization of linear functional
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# $E_{g} = \underset{\Gamma \in \mathcal{P}}{\operatorname{Min}} \operatorname{Tr} H\Gamma$ $\mathcal{P} = \{\Gamma : \operatorname{Approx} N \operatorname{-rep.condition}\}$

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P,Q,G,T1,T2-matrix are all positive semidefinite  $\leftrightarrow$  eigenvalues  $\lambda_i \ge 0$ 

$$U^{\dagger} \Gamma U = \begin{bmatrix} \lambda_1 & & 0 \\ & \lambda_2 & \\ & & \ddots & \\ 0 & & & \lambda_n \end{bmatrix} \ge 0$$

First application to Be atom [Garrod et al 1975, 1976] Calculation methods are not very well studied...

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[Nakata-Nakatsuji-Ehara-Fukuda-Nakata-Fujisawa 2001] [Nakata-Nakatsuji-Ehara 2002]

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Small enough "primal dual gap, feasibility" values show that total energies etc are MATHEMATICALLY correct

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- polynomial algorithm (cf. Hartree-Fock is NP-hard).

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- P, Q and G condition: 100 ~ 130% COrr. [Nakata et al], [Mazziotti et al] [Eric et al]
- P, Q, G, T1, T2' condition:  $100 \sim 101\%$  corr. [Zhao et al], [Nakata et al]
- *P*, *Q* and *G* condition: dissociation limit (sometimes fails).

[Nakata et al], [Mazziotti], [H. Aggelen et al]

#### The ground state energy of atoms and molecules [Nakata et al 2008]

System	1 State	N r	$\Delta E_{GT1T2}$	$\Delta E_{GT1T2'}$	$\Delta E_{CCSD(T)}  \Delta E_{HF} \qquad E_{FCI}$
С	<sup>3</sup> P	6 20	-0.0004	-0.0001	+0.00016 +0.05202 -37.73653
0	$^{1}D$	8 20	-0.0013	-0.0012	+0.00279 +0.10878 -74.78733
Ne	<sup>1</sup> S	10 20	-0.0002	-0.0001	-0.00005 +0.11645 -128.63881
0,+	$^{2}\Pi_{g}$	15 20	-0.0022	-0.0020	+0.00325 +0.17074 -148.79339
ВĤ	$^{1}\Sigma^{+}$	6 24	-0.0001	-0.0001	+0.00030 +0.07398 -25.18766
CH	$^{2}\Pi_{r}$	7 24	-0.0008	-0.0003	+0.00031 +0.07895 -38.33735
NH	$^{1}\Delta$	8 24	-0.0005	-0.0004	+0.00437 +0.11495 -54.96440
HF	$^{1}\Sigma^{+}$	14 24	-0.0003	-0.0003	+0.00032 +0.13834 -100.16031
SiH₄	${}^{1}A_{1}$	18 26	-0.0002	-0.0002	+0.00018 +0.07311 -290.28490
F-	<sup>1</sup> S	10 26	-0.0003	-0.0003	+0.00067 +0.15427 -99.59712
Р	<sup>4</sup> S	15 26	-0.0001	-0.0000	+0.00003 +0.01908 -340.70802
$H_2O$	${}^{1}A_{1}$	10 28	-0.0004	-0.0004	+0.00055 +0.14645 -76.15576

GT1T2	:	The RDM method (P, Q, G, T1 and T2 conditions)
GT1T2'	:	The RDM method ( $P, Q, G, T1$ and $T2'$ conditions)
CCSD(T)	:	Coupled cluster singles and doubles with perturbation treatment of triples
HF	:	Hartree-Fock
FCI	:	FullCI

#### Application to potential energy curve

 Dissociation curve of N<sub>2</sub> (triple bond) the world first result. [Nakata-Nakatsuji-Ehara 2002]





#### Recent results: non-size extensivity

Size extensivity or consistency is very important property for a calculation theory.

$$E(A - -infinity - -A) = E(A) + E(A)?$$

 Not size consistnt: [Nakata-Nakatsuji-Ehara 2002] (small deviation), [Aggelen-Bultinck-Verstichel-VanNeck-Ayers 2009] (fractional charge!)

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- Not size extensive: [Nakata-Yasuda 2009] PRA80,042109(2009).
  - CH<sub>4</sub>, N<sub>2</sub> non interacting polymers: slightly deviated
  - primal-dual interior point method is mandatory; Monteiro-Bruner [Mazziotti 04] is inaccurate.

#### Size-extensivity: N<sub>2</sub> polymer

 $N_2 N_2 N_2 \cdots N_2$  non interacting, *N*-rep.: *PQG* 



#### $E(M) = -108.71553 + 0.00302M^{-2}$ . $3 \times 10^{-4}$ au

#### Size-extensivity: CH<sub>4</sub> polymer

CH<sub>4</sub> CH<sub>4</sub> CH<sub>4</sub> ··· CH<sub>4</sub> non interacting, N-rep.: PQG



#### Nither PQG nor PQGT1T2' are size-extensive

#### Size-extensivity: Inaccurate result by Monteiro-Bruner method

 $H_2O$ : solved by Monteiro-Bruner method [Mazziotti 2004]: # of iteration req'ed scale like exponential. Not converged with CO (double- $\zeta$ ).


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# How many iterations are needed?

How many iterations are required by

- primal-dual interior-point method (PDIPM) or
- Monteiro-Bruner method (RRSDP) [Mazziotti 2004]

	P, Q, and G			P, Q, G, T1, T2		
algorithm	flops	# iterations	memory	flops	# iterations	memory
PDIPM	<i>r</i> <sup>12</sup>	$r \ln \varepsilon^{-1}$	$r^8$	r <sup>12</sup>	$r^{3/2}\lnarepsilon^{-1}$	$r^8$
RRSDP $r^6$ none $r^4$ $r^9$ none $r^6$						
Note: when we stop the iteration is a big problem						

### How large these SDP are?

# of constraints			
r	constraints	onstraints block	
24	15018	2520x2, 792x4, 288x1,220x2	
26	20709	3211x2, 1014x4, 338x1, 286x2	

Elapsed time using Itanium 2 (1.3GHz) 1 node 4 processors.

System, State, Basis	N-rep.	r	Time	# of nodes
SiH <sub>4</sub> , <sup>1</sup> A <sub>1</sub> , STO-6G	PQGT1T2	26	5.1 days	16
$\mathrm{H}_{2}\mathrm{O},^{1}A_{1},\mathrm{double}$ - $\zeta$	PQG	28	2.2 hours	8
$\mathrm{H}_{2}\mathrm{O},^{1}A_{1},\mathrm{double}$ - $\zeta$	PQGT1T2	28	20 days	8
$\mathrm{H}_{2}\mathrm{O},^{1}A_{1},\mathrm{double}$ - $\zeta$	PQGT1T2'	28	24 days	8

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- double (16 digits)  $1 + 0.0000000000000001 \simeq 1$
- GMP (GNU multiple precision) ⇒ necessity of highly accurate solver, using multiple precision arithmetic (SDPA-GMP) http://sdpa.indsys.chuo-u.ac.jp/sdpa/ GNU Public License

## SDPA-GMP and Hubbard model

The 1D Hubbard model with high correlation limit  $|U/t| \rightarrow \infty$ : All states are almost degenerated.

The ground state energies of 1D Hubbard model

PBC, # of sites:4, #	of electrons: 4, spin 0
----------------------	-------------------------

U/t	SDPA (16 digits)	SDPA-GMP (60 digits)	fullCI		
10000.0	0	$-1.1999998800000251 \times 10^{-3}$	$-1.199999880 \times 10^{-3}$		
1000.0	$-1.2 \times 10^{-2}$	$-1.1999880002507934 \times 10^{-2}$	$-1.1999880002 \times 10^{-2}$		
100.0	$-1.1991 \times 10^{-1}$	$-1.1988025013717993 \times 10^{-1}$	$-1.19880248946 \times 10^{-1}$		
10.0	-1.1000	-1.0999400441222934	-1.099877772750		
1.0	-3.3417	-3.3416748070259956	-3.340847617248		
PBC, # of sites:6, # of electrons: 6, spin 0					
U/t	SDPA (16 digits)	SDPA-GMP (60 digits)	fullCI		
10000.0	0	$-1.7249951195749525 \times 10^{-3}$	$-1.721110121 \times 10^{-3}$		
1000.0	$-1 \times 10^{-2}$	$-1.7255360310431304 \times 10^{-2}$	$-1.7211034713 \times 10^{-2}$		
100.0	$-1.730 \times 10^{-1}$	$-1.7302157140594339 \times 10^{-1}$	$-1.72043338097 \times 10^{-1}$		
10.0	-1.6954	-1.6953843276854447	-1.664362733287		
1.0	-6.6012	-6.6012042217806286	-6.601158293375		