# Piecewise polynomial approximation algorithm for short-range intermolecular interaction on wide SIMD architectures

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## **Background**

- Interaction in molecular dynamics simulations, which can divided into shortand long-range part in Ewald sum method, dominates the calculation time.
- To accelerate, the short-range force calculation has been replaced with rational approximation, table lookup interpolation, polynomial approximation.
- Piecewise polynomial approximation is a powerful technique to approximate functions with low order polynomials keeping accuracy for small sections.

Pro: multiply and add operation rich, division free.

Con: Random memory access to table of polynomial coefficients.

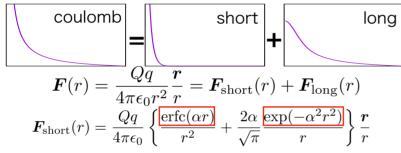
512-bit-width SIMD registers of novel high-performance CPUs can serve as an efficient table for piecewise polynomial approximation.

# **Objective**

To demonstrate a new implementation of piecewise polynomial approximation algorithm using wide SIMD registers as a table to accelerate the calculation of short-range intermolecular interaction.

# **Algorithm**

#### Short-range interaction in Ewald sum method



Q,q: charges, r: distance between charges,  $\varepsilon_0$ : permittivity of vacuum

Coulomb interaction is divided into two part, short- and long-range part. The short-range part includes complementary error function and exponential function which results in highly compute intensive.

#### Existing method: Rational function approximation (RFA)

## GROMACS\*, one of the fastest MD simulation package, adopts RFA.

 $m{F}_{
m short}(r) = rac{Qq}{4\pi\epsilon_0} \left\{ rac{1}{r^3} - rac{{
m erf}(lpha r)}{r^3} + rac{2lpha}{\sqrt{\pi}} rac{{
m exp}(-lpha^2 r^2)}{r^2} 
ight\} m{r}$   $\sum_{m=0}^{10} b_m (lpha^2 r^2)^m \over \sum_{m=0}^{5} c_m (lpha^2 r^2)^m} b_m b_m (eta), c_m (eta)$ : mth coefficient of polynomial function

The correlation part of short-range force is approximated by rational function. For details, see pmeForceCorrectionSingleAccuracy on http://manual.gromacs.org/documentation/current/doxygen/html-lib/group\_module\_simd.xhtml.

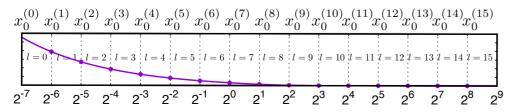
\*Abraham, et al., Software X, 1-2, 2015.

#### New method:

### Piecewise polynomial approximation (PPA)

$$F_{\rm short}(r) = \frac{Qq}{4\pi\epsilon_0} \left\{ \begin{array}{l} {\rm erfc}(\sqrt{(\alpha r)^2}) + \frac{2}{\sqrt{\pi}} \exp(-\alpha^2 r^2) \sqrt{(\alpha r)^2} \right\} \frac{r}{r^3} \\ x = (\alpha r)^2 - x_0^{(l)} \\ F_{\rm approx}^{(l)}(x) = \left( \sum_{m=0}^n a_m^{(l)} x^m \right)^2 \frac{\alpha : {\rm constant\ of\ Ewald\ sum\ method.}}{x_0^{(l)} : {\rm lower\ bound\ of\ } (\alpha r)^2 {\rm\ in\ region\ } l.} \\ a_m^{(l)} : {\rm\ mth\ coefficient\ of\ polynomial\ function\ in\ region\ } l. \end{array} \right.$$

Deformation of original short-range force formula to make target function not change drastically brings higher accuracy.



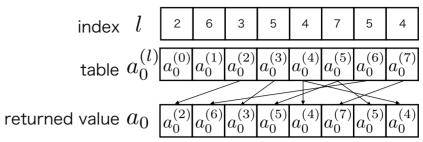
To make approximated functions accurate,

- Intervals  $(x_0^{(l)})$  are set to power of 2.
- Sollya\* is used for computing minimax polynomials of each 1.

# \*S. Chevillard, M. Joldeş and C. Lauter. "Sollya: an environment for the development of numerical codes," In *Mathematical Software - ICMS 2010*, pages 28-31, Heidelberg, Germany, September 2010. Springer.

# **Implementation**

Table lookup/permute instruction for 512-bit SIMD



512-bit SIMD registers can be used as a table with instructions below: For Intel architecture

- \_\_m512 \_mm512\_permutexvar\_ps(\_\_m512i idx, \_\_m512 a)
- \_m512 \_mm512\_permutex2var\_ps(\_m512 a,\_\_m512i idx, \_\_m512 b)
   For ARM v8 architecture
- svfloat32\_t svtbl\_f32(svfloat32\_t a, svuint32\_t idx)

#### Pseudo code of PPA

#### Require:

 $a_m^{(l)}$ : polynomial coefficients.

X.f: X as an floating point value w/o changing any bit.

X.s : X as an integer value w/o changing any bit.

offset: offset binary of exponent bit.

#### function *n*th-polynomial approximation {

$$r^{2} = |r_{i} - r_{j}|^{2};$$

$$R^{2} = \alpha^{2} * r^{2};$$

$$l = (((R^{2}).s - offset) >> 23) & 0x0f;$$

$$x_{0}^{(l)} = ((R^{2}).s & 0xff800000).f;$$

$$x = R^{2} - x_{0}^{(l)};$$

$$return \left(\sum_{m=0}^{n} a_{m}^{(l)} x^{m}\right)^{2};$$

## Accuracy

Comparison with the original force in DP and PPA in SP.

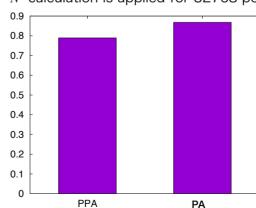
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# Benchmark result

N<sup>2</sup> calculation is applied for 32768 point charge distributed on cubic lattice.



CPU: Intel Xeon Gold 6140@2.30GHz
Compiler: icc 17.0.0 20160721
Option: -xCORE-AVX512 -ip
-unroll-loops
-use-fast-math

r / Angstrom

PPA is 10% faster than the previously known polynomial-approximation implementation.

## **Conclusion**

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Elapsed

We proposed a piecewise polynomial approximation algorithm which uses SIMD registers as a table of polynomial coefficients for efficient calculation of short-range intermolecular interaction.

- Enough accurate for for the calculation in single-precision.
- 10% faster than the best previously known SIMD implementation on Intel Skylake Xeon.