

Accounting for cross-association in nonself-associating species using a physically consistent SAFT-VR Mie approach J.T. Cripwell*, S.A.M. Smith, C.E. Schwarz, A.J. Burger

Department of Chemical Engineering, Stellenbosch University, Banghoek Road, Stellenbosch 7600, South Africa. Email: *cripwell@sun.ac.za. Tel: +27 21 808 4108

BACKGROUND

- Self-associating (SA) molecules have both electron pair donor and acceptor sites, and can form hydrogen bonds with themselves
- Molecules with only donor- or only acceptor sites, such as ethers, ketones, and esters, are nonself-associating (NSA), but can form hydrogen bonds with other molecules → cross-association
- Traditional SAFT parameterisation only treats SA molecules explicitly
- Need approach to obtain association parameters for NSA molecules

mole fraction 2-butanone

Fig. 2. 2-Butanone + ethanol VLE at three temperatures. Data from Ref. [7].

mole fraction 2-butanone

• Results indicate temperature dependence of association parameters

To develop a general and physically consistent approach to account for cross-association in a predictive manner for a wide range of components

METHODOLOGY

STEP 1: Select thermodynamic model

- SAFT-VR Mie-GV [1] combines SAFT-VR Mie [2] with the dipolar term of Gross & Vrabec [3]
- SAFT-VR Mie-GV accurately describes properties of dipolar molecules, therefore serves as suitable foundation of this work
- A single electron pair donor site (**N scheme**) is assigned to the polar, NSA molecules that will allow them to cross-associate [4,5]

STEP 2: Select tool to determine association parameters

- **Discretised sensitivity analysis** [4,5] is used to determine suitable association parameters
- The SAFT-VR Mie-GV parameters are fixed, and the association parameters (ε^{AB} and r_c) are varied at discrete intervals
 Mixture VLE is calculated at each combination of (ε^{AB}, r_c)
 The deviations are projected onto the (ε^{AB}, r_c)-plane, resulting in contour plots (see Fig. 1(a))
 In this work an "alcohol-first" approach is followed: binary VLE of alcohol + ether/ketone/ester systems are used in the development

• Highlights potential shortcoming of association term

2. Molecules not included in method development



Fig. 3. (a) Isopropyl ethanoate + 2-propanol [8]; (b) Pentyl ethanoate + 1-pentanol [9].

• Application *beyond* components used in the method development support validity of approach and correlation used for esters

3. Aqueous mixtures

mole fraction 2-butanone



STEP 3: Analysing discretisation results

- $(\varepsilon^{AB}, r_{c})$ -pairs on the same contour give almost identical predictions
- Ketone/ether mixtures: minimum AAD locus (white line in Fig. 1(a)) is similarly located for all investigated systems → can use *global* parameter set for homologous group
- For ester mixtures, $\varepsilon^{AB} = f(r_c^{\text{alcohol}}, M_W^{\text{ester}})$



Fig. 4. (a) Acetone + water VLE [10]; (b) Methyl ethanoate + water LLE [11,12].

- Two negative sites (2N) required for qualitative VLE description
- LLE description lacking, even with inclusion of k_{ij}

4. Mixtures with chloroform



Fig. 5. (a) Acetone + chloroform [13]; (b) Ethyl ethanoate + chloroform [14].

Fig. 1. Contour plot (a) and corresponding predictions (b) for methyl ethanoate +

ethanol. Data from Ref. [6].

• Obtain correct qualitative description using P-scheme for chloroform

CONCLUSIONS

- Cross-association offers predictive approach for solvating systems
- Approach can extrapolate to components not involved in development
- Temperature dependence of association contribution evident
- Description of aqueous mixtures remains difficult

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