

XMLTDB in FactSage Calphad Optimizer and ChemApp: Lessons learned and opportunities generated

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Current XMLTDB proposal (0.0.6)

```
<Database version="0.0.6">
  <metadata>
    <writer Software="OpenCalphad 6.068" Date="2023-10-26" />
  </metadata>
  </metadata>
  <Defaults LowT="10" HighT="6000" Bibref="U.N. Known" Elements="VA /-"/>
  <Element Id="AL" Refstate="FCC_A1" Mass="26.982" H298="4577.3" S298="28.322" />
  <Element Id="C" Refstate="GRAPHITE" Mass="12.011" H298="1054" S298="5.7423E+00" />
  <Species Id="VA" Stoichiometry="VA" />
  <Species Id="AL" Stoichiometry="AL" />
  <Species Id="C" Stoichiometry="C" />
  <TPfun Id="R" Expr="8.31451;" />
  <TPfun Id="RTLNS" Expr="R*T*LN(1.0E-5*P);" />
  <TPfun Id="GOAL4C3" Expr="-277339-.005423368*T**2;" />
  <TPfun Id="GTSERCC" Expr="-0.01478307*T**2-7.83339395E-07*T**3;" />
  <TPfun Id="GOBCCAL" Expr="-0.0029531333*T**2-3.3998492E-16*T**5;" />
  <TPfun Id="GOCPCAL" Expr="+GHSERAL+10083;" />
  <TPfun Id="GOCPCAL" Expr="+GHSERAL+5481;" />
  <TPfun Id="GHSERCC" Expr="-8160*GTSERAL;" />
  <TPfun Id="GHSERCC" Expr="-17752.213+GGRACCC+GTSERCC;" />
  <TPfun Id="GODIACC" Expr="-16275.202-9.1299452E-05*T**2-2.1653414E-16*T**5;" />
  <TPfun Id="GEDIACC" Expr="+0.2318*GEIN(+813.6)+.01148*GEIN(+345.3)-0.236743*GEIN(1601);" />
  <TPfun Id="GOLIQAL" Expr="-209-3.777*T-.00045*T**2;" />
  <TPfun Id="GOLIQCC" Expr="+63887-8.2*T-.0004185*T**2;" />
  <TPfun Id="GGRACCC" Expr="-0.5159523*GEIN(+1953)+0.121519*GEIN(+448)+0.3496843*GEIN(+9411)+0.0388463*GEIN(+192.65)+.005840323*GEIN(+64.46);" />
  <Phase Id="LIQUID" Configuration="CEF" State="L">
    <Sites NumberOf="1" Multiplicities="1">
      <Constituents Sublattice="1" List="AL C" />
    </Sites>
    <AmendPhase Models="LIQSTATE" />
  </Phase>
  <Phase Id="AL4C3" Configuration="CEF" State="S">
    <Sites NumberOf="2" Multiplicities="4 3">
      <Constituents Sublattice="1" List="AL" />
      <Constituents Sublattice="2" List="C" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Phase Id="BCC_A2" Configuration="CEF" State="S">
    <Sites NumberOf="2" Multiplicities="1 3">
      <Constituents Sublattice="1" List="AL" />
      <Constituents Sublattice="2" List="C VA" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Phase Id="DIAMOND" Configuration="CEF" State="S">
    <Sites NumberOf="1" Multiplicities="1">
      <Constituents Sublattice="1" List="C" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Phase Id="FCC_A1" Configuration="CEF" State="S">
    <Sites NumberOf="2" Multiplicities="1 1">
      <Constituents Sublattice="1" List="AL" />
      <Constituents Sublattice="2" List="C" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Phase Id="GRAPHITE" Configuration="CEF" State="S">
    <Sites NumberOf="1" Multiplicities="1">
      <Constituents Sublattice="1" List="C" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Phase Id="HCP_A3" Configuration="CEF" State="S">
    <Sites NumberOf="2" Multiplicities="1 0.5">
      <Constituents Sublattice="1" List="AL" />
      <Constituents Sublattice="2" List="C VA" />
    </Sites>
    <AmendPhase Models="GEIN" />
  </Phase>
  <Parameter Id="G(LIQUID,AL;0)" Expr="+GOLIQAL" Bibref="20HE" />
  <Parameter Id="LNTH(LIQUID,AL;0)" Expr="+LN(1400);" Bibref="20HE" />
  <Parameter Id="G2(LIQUID,AL;0)" Expr="+13398-R*T-0.169" Bibref="20HE" />
  <Parameter Id="G(LIQUID,C;0)" Expr="+GOLIQCC;" Bibref="20HE" />
  <Parameter Id="LNTH(LIQUID,C;0)" Expr="+LN(1400);" Bibref="20HE" />
  <Parameter Id="G2(LIQUID,AL,C;0)" Expr="+59147-49.61*T+2.9806*T*LN(T);" Bibref="20HE" />
  <Parameter Id="G(LIQUID,AL,C;0)" Expr="+20994-22*T;" Bibref="20HE" />
  <Parameter Id="G(AL4C3,AL;C;0)" Expr="+GOAL4C3-3.08*GEIN(+401)+3.08*GEIN(+1077);" Bibref="20HE" />
  <Parameter Id="LNTH(AL4C3,AL;C;0)" Expr="+LN(401);" Bibref="20HE" />
  <Parameter Id="G(BCC_A2,AL;C;0)" Expr="+GTSERAL+3*GTSERCC+1006844;" Bibref="20HE" />
  <Parameter Id="LNTH(BCC_A2,AL;C;0)" Expr="+LN(863);" Bibref="20HE" />
  <Parameter Id="G(BCC_A2,AL;VA;0)" Expr="+GOBCCAL;" Bibref="20HE" />
  <Parameter Id="LNTH(BCC_A2,AL;VA;0)" Expr="+LN(233);" Bibref="20HE" />
  <Parameter Id="G(BCC_A2,AL,C,VA;0)" Expr="-819896+14*T;" Bibref="20HE" />
  <Parameter Id="G(DIAMOND,C;0)" Expr="+GODIACC+GEDIACC;" Bibref="20HE" />
  <Parameter Id="LNTH(DIAMOND,C;0)" Expr="+LN(1601);" Bibref="20HE" />
  <Parameter Id="G(FCC_A1,AL;C;0)" Expr="+GTSERAL+GTSERCC;" Bibref="20HE" />
  <Parameter Id="LNTH(FCC_A1,AL;C;0)" Expr="+LN(549);" Bibref="20HE" />
  <Parameter Id="G(FCC_A1,AL;VA;0)" Expr="+GHSERAL;" Bibref="20HE" />
  <Parameter Id="LNTH(FCC_A1,AL;VA;0)" Expr="+LN(283);" Bibref="20HE" />
  <Parameter Id="G(FCC_A1,AL,C,VA;0)" Expr="-70345;" Bibref="20HE" />
  <Parameter Id="G(GRAPHITE,C;0)" Expr="+GHSERCC;" Bibref="20HE" />
  <Parameter Id="LNTH(GRAPHITE,C;0)" Expr="+LN(1953);" Bibref="20HE" />
  <Parameter Id="G(HCP_A3,AL;C;0)" Expr="+GTSERAL+0.5*GTSERCC+2176775;" Bibref="20HE" />
  <Parameter Id="LNTH(HCP_A3,AL;C;0)" Expr="+LN(452);" Bibref="20HE" />
  <Parameter Id="G(HCP_A3,AL;VA;0)" Expr="+GOCPCAL;" Bibref="20HE" />
  <Parameter Id="LNTH(HCP_A3,AL;VA;0)" Expr="+LN(263);" Bibref="20HE" />
  <Parameter Id="G(HCP_A3,AL,C,VA;0)" Expr="0;" Bibref="20HE" />
  </Bibliography>
  <Bibliography Id="20HE" Text="Zhang, Tang, He, Sundman, Calphad" />
</Database>
```

Defaults are comfortable, but can be dangerous, e.g. when combining data sets

Very closely modeled after PHASE and CONSTITUENTS commands in TDBs

Model description is nested in phase tag – where are required model parameters?

Other models: MQMA, I2SL, UNIQUAC, ...
What is a configuration model?
E.g. Redlich-Kister, Toop (more?)

A lot of information 'coded' into a small data field that is computationally 'hard' to decode

Definition and location of GTSERAL and GTSERCC unclear

All parameter lines can be put anywhere in the file.
Convention is: 'by system'

Suggestions and potential future features

Take advantage of XML properties

- All parameters collected under phase
- Parameters don't 'float' freely in file
- Schema validation can verify structural correctness

```
<Phase Id="BCC_A2" Configuration="CEF" State="S">
  <Sites NumberOf="2" Multiplicities="1 3">
    <Constituents Sublattice="1" List="AL" />
    <Constituents Sublattice="2" List="C VA" />
  </Sites>
  <AmendPhase Models="GEIN" />
  <Parameter MPID="G" ConstArray="AL:VA" Expr="+GOBCCAL;" Bibref="20HE" />
  <Parameter MPID="LNTH" ConstArray="AL:VA" Expr="+LN(233);" Bibref="20HE" />
  <!-- Al-C binary -->
  <Parameter MPID="G" ConstArray="AL:C" Expr="+GTSERAL+3*GTSERCC+1006844;" Bibref="20HE" />
  <Parameter MPID="LNTH" ConstArray="AL:C" Expr="+LN(863);" Bibref="20HE" />
  <Parameter MPID="L" ConstArray="AL:C,VA" Degree="0" Expr="-819896+14*T;" Bibref="20HE" />
</Phase>
```

As it is nested in the BCC_A2 phase tag, the MPID can omit the phase id

separating MPID makes this more explicit

Unification of 'primitive' technical requirements

- Phase name lengths: 25 characters? 28 characters? Unlimited?
- (dis)allowed characters
- Can end members have 'common names'?
- Number precision defaults, esp. for output?
- Prevent circular function references

Modern world requirements: FAIR data support

- Verification: checksums, cryptographic signing
- Change management:
 - 'beautiful diff' for versioning
 - tracking changes and maintain evolution
- Ontology adaption (integrate into larger data concepts)
- Comments and other meta-data on data/values

Connect atomistic model to phase model

- Allows description of first principle models
- Synchronization with e.g. VASP input / output

```
<Phase Id="HypotheticalMu" Configuration="CEF" State="S">
  <AmendPhase Models="ReuterCryst Vegard" />
  <Sublattices NumberOf="3" Multiplicities="7 2 4">
    <Constituents Sublattice="1" List="CO CR FE NI TA HF" />
    <Constituents Sublattice="2" List="MO NB TA W HP" />
    <Constituents Sublattice="3" List="CO CR FE MO NB NI TA W HP" />
  </Sublattices>
  <Crystallography>
    <SpaceGroup number="166">
      <!-- General configuration -->
      <Configuration const="*:*:*">
        <Wyckoff Symbol="3b" Sublattice="1" />
        <Wyckoff Symbol="18h" x="0.5000" z="0.0900" Sublattice="1" />
        <Wyckoff Symbol="6c1" z="0.0520" Sublattice="2" />
        <Wyckoff Symbol="6c2" z="0.1540" Sublattice="3" />
        <Wyckoff Symbol="6c3" z="0.3333" Sublattice="3" />
        <Parameter MPID="LP" a="4.764" b="4.764" c="25.850" Bibref="" />
        <Parameter MPID="LA" a="90" b="90" c="120" Bibref="" />
      </Configuration>
      <!-- Specific to one endmember -->
      <Configuration const="CO:N:FE" Name="Hypothetical Special Case">
        <Wyckoff Symbol="3b" Sublattice="1" />
        <Wyckoff Symbol="18h" x="0.5001" z="0.0900" Sublattice="1" />
        <Wyckoff Symbol="6c1" z="0.0521" Sublattice="2" />
        <Wyckoff Symbol="6c2" z="0.1541" Sublattice="3" />
        <Wyckoff Symbol="6c3" z="0.3334" Sublattice="3" />
        <Parameter MPID="LP" a="4.765" b="4.765" c="25.851" Bibref="" />
        <Parameter MPID="LA" a="90" b="90" c="120" Bibref="" />
      </Configuration>
    </SpaceGroup>
  </Crystallography>
</Phase>
```

Key crystallographic information to conceptualize the phase model

Relation of Wyckoff sites and sublattice modeling explicitly

Current status (for proposal v0.0.6):

- Reading/Writing from OpenCalphad
- Reading/Writing from ChemApp
- Schema validation XSL file available

Roadmap:

- Version 1.0.0 by the end of the year
- Getting Feedback!

Florian Tang

Contact info



<https://github.com/flotang-gtt/XMLTDB>



<https://github.com/sundmanbo/XMLTDB>



Biggest questions

- Uncertainty quantification*
If a dataset has been researched and evaluated, how can this information be attached to the model?
How will this data be supported in the software tools?
- Stewardship for XMLTDB*
Is it necessary to have authority over the format?
- Beyond current database data*
 - Safety margins, assessed modeling ranges
 - More meta data – e.g. warnings, dependencies

