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1 **Yttrium complexation and hydration in chloride-rich hydrothermal**
2 **fluids: a combined *ab initio* molecular dynamics and *in situ* X-ray**
3 **absorption spectroscopy study**

4 Qiushi Guan^{1,2}, Yuan Mei^{2*}, Barbara Etschmann¹, Denis Testemale³, Marion Louvel⁴,
5 Joël Brugger^{1*}

6 1. School of Earth, Atmosphere and the Environment, Monash University,
7 Clayton, VIC 3800, Australia

8 2. CSIRO Mineral Resources, Kensington WA 6152, Australia

9 3. CNRS, Université Grenoble Alpes, Institut NEEL, F-38000 Grenoble, France

10 4. Institute for Mineralogy, WWU Muenster, D-48149 Germany

11

12 *Corresponding author: Yuan Mei (Yuan.Mei@csiro.au); Joël Brugger
13 (joel.brugger@monash.edu)

14

15 **Abstract**

16 Accurate knowledge of rare earth elements (REE) speciation in high pressure –
17 high temperature fluids is required to model REE transport and precipitation in
18 subduction zones and magmatic-hydrothermal environments, and the formation of
19 rare metal deposits. Recent experiments (lanthanum, ytterbium, erbium) have
20 demonstrated that REE chloride complexes are the main REE form in many
21 hydrothermal fluids (Migdisov, et al., 2016). However, the speciation of yttrium

22 (Y(III)), a cation with an ionic radius similar to that of Ho(III), remains poorly
23 constrained in chloride-rich hydrothermal solutions.

24 We used *ab initio* molecular dynamics (MD) simulations to calculate the nature
25 of Y(III)-Cl complexes and the thermodynamic properties of these species at
26 temperatures up to 500 °C and pressures of 800 bar and 1000 bar. The MD results
27 were complemented by *in-situ* X-ray absorption spectroscopy (XAS) measurements.
28 Our results indicate that at temperature below 200 °C, chloro-complexes do not form
29 readily, even in highly concentrated brines. At ambient condition, the Y(III) aqua ion
30 binds to eight water molecules in a square antiprism geometry, which is consistent
31 with previous *ab initio* studies (Ikeda, et al., 2005). The thermodynamic integration
32 method was employed to calculate the formation constants ($\log\beta$) of Y(III)-Cl⁻
33 complexes in two simulation boxes containing different Y:Cl ratios; we obtained very
34 consistent results of the standard $\log\beta$ of same complexes from the two independent
35 calculations, which confirms that the thermodynamic integration method is reliable
36 and not significantly affected by technical limitations in box size, box composition, or
37 simulation times.

38 Based on the derived formation constants, we fit Modified Ryzhenko–Bryzgalin
39 (MRB) equation of state parameters, which enable extrapolation of the formation
40 constants at elevated temperature and pressure. The results are consistent with the
41 XAS data, and show that the stability of Y(III)-Cl complexes increases with
42 increasing temperature, Y(III) forming high order Cl⁻ complexes (up to YCl₄⁻) in high

43 salinity solutions at high temperature and $\text{pH} = 3$. We also compare the extrapolated
44 $\log\beta$ with the available data for other REE at 150 °C, 200 °C and 250 °C. At 200 °C,
45 yttrium behaves more like a heavy REE, but from 200 °C to 250 °C, the $\log\beta$ of
46 Y(III)-Cl complexes increases dramatically and behave more like the light REE. The
47 difference of Cl^- dominant species between Ho(III) (HoCl_2^{2+}) and Y(III) (YCl_2^+) may
48 account for the formation of anomalous Y/Ho ratios in some hydrothermal
49 environments.

50

51 Keywords: Yttrium; *ab initio* molecular dynamics; XAS; yttrium chloride aqueous
52 speciation; thermodynamics; rare earth elements.

53

54

55

56 **1. Introduction**

57 Demand for rare earth elements (REE: La to Lu) plus Y has increased in the past
58 decades due to their use in fast-growing industrial applications (e.g., energy, mobility
59 and special alloys; [Krishnamurthy, 2016](#)). Combined with strategic issues related to
60 current concentration of REE production within China, this has led to world-wide
61 efforts to secure new geological sources of REE ([Weng et al., 2015](#); [Ram et al. 2018](#)).
62 The world's largest REE deposit formed by a combination of magmatic and
63 hydrothermal processes (Bayan Obo, China; [Ling et al., 2013](#)). REE are mobile in
64 many hydrothermal systems (e.g., Iron Oxide Copper Gold deposits and their
65 alteration halos; ([Kontonikas-Charos et al., 2018](#); [Xing et al., 2019](#)), and since REE
66 behave coherently through geochemical processes, they can be used as tracers for
67 fluid-rock interaction (e.g., [Bau and Dulski, 1995](#); [Brugger et al., 2008](#)). Over the past
68 decade, a number of experimental studies have dramatically improved our
69 understanding of REE speciation in natural water at elevated temperature, e.g. *in situ*
70 measurements at high temperature-pressure for Yb ([Louvel et al. 2015](#)) and Eu ([Liu et](#)
71 [al. 2017](#)); solubility experiments of REE+Y-fluoride ([Loges et al. 2013](#); [Migdisov](#)
72 [2009](#)); *in situ* spectrophotometric studies of REE chloride and sulfate complexes
73 ([Migdisov and Williams-Jones, 2008](#); [Migdisov et al. 2008](#)). As reviewed by
74 [Migdisov et al. \(2016\)](#), the nature of REE complexes (e.g., chloride versus fluoride or
75 hydroxide complexes) affects mineral solubility, and is controlled by temperature,
76 pressure and solution composition.

77 A molecular-level understanding of the coordination geometry and hydration of
78 the complexes is key to reliable extrapolations of thermodynamic properties that can
79 be also used to model metal mobility beyond the conditions covered by experiments
80 (Brugger et al., 2016). Chloride is the most common ligand in many geo-fluids, and
81 despite the fact that REE fluoride complexes are much stronger than chloride
82 complexes (Haas et al., 1995; Wood, 1990), chloride complexes are considered as the
83 main transporting ligands in many hydrothermal fluids, especially at high temperature,
84 since the stability of chloride complexes increases with increasing temperature
85 relative to fluoride complexes, and the activity of the fluoride ion is limited by the
86 strong association of hydrofluoric acid at elevated temperature and the low solubility
87 of fluorine-containing phases such as fluorite, apatite, and biotite (Gammons et al.,
88 1996; Migdisov et al., 2009; Xing et al. 2019).

89 The similar chemical properties of REE *sensu stricto* (lanthanides) are a result of
90 their valence electrons in $4f$ -orbitals; these f -electrons are buried deep within the atom
91 and shielded by $4d$ and $5p$ electrons and do not take part in bonding. Yttrium is often
92 grouped with REE because its ionic radius (0.900 Å for Y^{3+}) is close to that of the
93 heavy REE (Ho^{3+} at 0.901 Å; Shannon, 1976). However, due to its lack of f -electrons
94 ($[Kr] 4d^1 5s^2$), Y(III) shows significant differences in behavior compared to REE
95 *sensu stricto*. For example, Loges (2013) showed that at temperatures below 250 °C,
96 the dominant Y(III) fluoride complex is YF_2^+ , compared to $REEF_2^+$ (HoF_2^+). These
97 differences can result in contrasting mobility of Y and REE in hydrothermal systems

98 (Williams-Jones, 2009), leading to the concept of ‘yttrium anomaly’ (Bau and Dulski,
99 1995).

100 Several experiments have been conducted on the hydration of Y(III) at low
101 temperature (Table 1). At room temperature, the Y(III) aqua ion is generally
102 recognized to have a coordination of 8 and a square antiprism structure (references in
103 Table 1). The early *in-situ* XAS study of Ragnarsdottir et al. (1998) provides the only
104 experimental data on Y(III) hydration under hydrothermal conditions. Ragnarsdottir et
105 al. (1998) obtained a higher than expected Y(III) hydration number at room
106 temperature (9.1 – 10.3), and showed a small decrease in hydration number to 8 at
107 temperature of 250 °C, with no further dehydration observed upon heating to 340 °C.

108 There are only a few studies about Y(III) chloride complexing, and again
109 Ragnarsdottir et al. (1998) is the only experimental study extending to hydrothermal
110 conditions (Table 1). Conclusions regarding the extent of Y-Cl complexation differ.
111 On one hand, Allen et al. (2000)’s EXAFS study found no evidence for Cl⁻ bonded
112 with Y(III) at ambient temperature, even in brines containing up to 14 m Cl⁻.
113 Ragnarsdottir et al. (1998) similarly found no evidence of Y-Cl complexing in a 0.1 m
114 YCl₃ solution to 340 °C, and in a 0.1 m YCl₃ + 0.23 m HCl + 2 m NaCl solution to
115 250 °C (Cl_{tot} = 2.53 m). On the other hand, Luo and Byrne (2001) report a stability
116 constant of 0.65 ± 0.05 for the formation of YCl²⁺ (reaction Y³⁺ + Cl⁻ = YCl²⁺) at
117 25 °C and zero ionic strength, and concluded that Y³⁺ formed weak complexes with
118 Cl⁻. Combining neutron scattering and EXAFS spectroscopy, Bowron and

119 Diaz-Moreno (2007) also support the formation of Y-Cl complexes in 1 m YCl₃
120 solution at room temperature, with a hydration number of 7.4 ± 0.5 in the distance
121 between 2.1 Å and 3.3 Å, and 0.8 ± 0.2 Cl⁻ between 2.5 Å and 3.5 Å, from the radial
122 distribution functions of Y-O and Y-Cl. Rudolph and Irmer (2015) concluded in a
123 recent Raman study that Y(III) and Cl⁻ can form weak chloro-complexes at room
124 temperature in YCl₃+HCl solutions (2.03 M YCl₃ in 1.0 M and 4 M HCl). They
125 propose that Y exists in 8-fold coordination as [Y(H₂O)_{8-n}Cl_n]⁺³⁻ⁿ in these solutions,
126 with chloride numbers $n \leq 2$. Stefanski and Jahn (2019) studied Y(III) speciation in
127 chloride- and fluoride-rich brines under subduction conditions (800 °C, 13/45 kbar)
128 with *ab initio* molecular dynamics, and concluded that chloride complexes
129 [Y(H₂O)₂₋₇Cl₁₋₅]^{+3-n(Cl)} are important species of Y mass transfer in subduction zones.

130 Table 1. Structure of aqueous Y(III) in literature

131 In this study, we combined *ab initio* molecular dynamics (MD) simulations and
132 *in situ* XAS measurements to quantify the nature, geometry, and stability of Y(III)
133 complexes in chlorine-rich fluids from room temperature to 500 °C, 1000 bar. In
134 particular, the formation constants of Y(III)-Cl complexes were calculated from *ab*
135 *initio* MD using the thermodynamic integration method. Our previous studies have
136 demonstrated that the combination of *ab initio* computational chemistry and *in situ*
137 spectroscopic experiments is a powerful approach to determining element speciation
138 and geometry in hydrothermal fluids (Mei et al., 2015a,b, 2016; Brugger et al., 2016;
139 Etschmann et al., 2018, 2019), providing accurate speciation data that can be used to

140 provide extrapolations to conditions beyond experimental scope.

141 **2. Method**

142 *2.1 Ab initio* molecular dynamics

143 The *ab initio* MD simulations were conducted with Car-Parrinello Molecular
144 Dynamics (CPMD, version 3.1.17, Car and Parrinello, 1985). In CPMD, the electrons
145 were treated as plane wave basis sets using density functional theory. The interactions
146 between valence electrons were depicted by BLYP exchange correlation functional
147 (Becke, 1988; Lee et al., 1988), which produces a good description of the structural
148 properties of water including O–O interaction, angular distributions, coordination
149 numbers, H-bond statistics, and radial distribution functions when compared with
150 experiments (Gillan et al., 2016; Lin et al., 2012). However, most generalized gradient
151 approximation functionals, including BLYP, tend to overestimate the ordering of the
152 structure of liquid water at ambient condition (e.g., Sprik, et al. 1996, Mei et al.,
153 2018). Bankura et al. (2014) showed that BLYP best reproduces the properties of
154 water at ambient conditions when using an elevated temperature for the simulations,
155 and it could be argued that there is a temperature offset of approximately 55°C
156 between the computed data and experiment in terms of solvent properties. Despite
157 these limitations, many previous studies obtained accurate results of aqueous
158 metal-ligand complexations at both room and high temperature using the BLYP
159 functional (e.g. Mei et al., 2015b; Stefanski and Jahn, 2019). A timestep of 3 a.u.
160 (0.073 fs) and fictitious mass of 400 a.u. to all atoms were employed in the simulation.

161 We used the NVT ensemble to conduct the simulations. The [density of the solution](#)
162 was chosen [according](#) to the equation of state of NaCl solution at similar chloride
163 concentration at targeted T and P ([Driesner, 2007](#); [Driesner and Heinrich, 2007](#)). The
164 temperature was fixed with Nosé-Hoover chain thermostat for ions and electrons.
165 Periodic boundary condition was applied to eliminate surface effects.

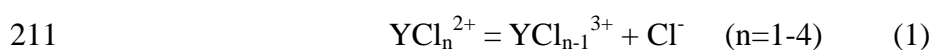
166 In total, 15 *ab initio* MD simulations were conducted over a range of
167 temperatures (25-500 °C), pressures (1, 800 and 1000 bar), and compositions
168 (simulation boxes containing 1 yttrium(III) ion, 3 to 5 chloride ion, and 55 or 110
169 H₂O molecules) relevant for a wide range of hydrothermal systems; the simulation
170 details are listed in [Table 2](#). All the MD simulations were conducted for 16-30
171 picoseconds (ps) in order to achieve good statistics. [As discussed in our previous](#)
172 [study \(Mei et al., 2013a\)](#), the size of the simulation boxes and times in this study can
173 [provide solid results within manageable computation resources](#). The time-averaged
174 stoichiometric and geometric information was obtained with VMD ([Humphrey et al.,](#)
175 [1996](#)). The coordination numbers were retrieved from radial distribution functions
176 (RDF), and their errors were estimated from the change of RDF integral at the
177 distance range of 2.75 Å - 3.25 Å for Y-O, and 2.9 Å - 3.2 Å for Y-Cl. The
178 Debye-Waller factor were calculated following equations in section 2.3 in [Campbell](#)
179 [et al. \(1999\)](#).

180 [Table 2. simulation details of ab initio MD](#)

181 2.2 *Ab initio* thermodynamic integration

182 The *ab initio* MD results reveal the structure of Y(III)-Cl complexes at different
183 P-T. However, *ab initio* MD is very resource-intensive: for example, obtaining
184 1 picosecond of real time data requires ~6,800 CPU hours for a simulation box
185 containing 111 waters with density of 0.73 g/cm³ at 450 °C, and ~1,300 CPU hours
186 for a simulation box containing 55 waters with density of 0.90 g/cm³ at 350 °C. This
187 limits the timescale of MD simulations to 10's of picoseconds (ps). Under these
188 conditions, observation of the distribution of species (i.e., statistical appraisal of
189 association/dissociation reactions of ligands) in a solution can be beyond current
190 computational capabilities, since ligand exchange rates are in the timescale of
191 nanoseconds or microseconds for many aqua- and chloride complexes. For example,
192 the water residence time of Ho³⁺ is 4.67 ns at 298 K (Cossy et al., 1989), with Y³⁺
193 expected to have a value within the same order of magnitude based on similarities in
194 the ionic radius of Y(III) and Ho(III) (Shannon, 1979). The residence times of the Cl⁻
195 ligand in complexes of the transition metals Au(III), Zn(II) and Cd(II) are around
196 9.3 ms, 251 μs and <32 μs, respectively (Sharps et al., 1993). Even though ligand
197 residence times decrease with temperature to a point where individual ligand
198 exchange reactions can be easily observed via *ab initio* MD, the exchange rates are
199 generally not sufficient to allow measurements of the relative complexes stabilities
200 from the distribution of species in the simulations. Instead, the thermodynamic
201 integration method (Sprik, 1998; Sprik & Ciccotti, 1998) was used to obtain

202 quantitative thermodynamic information on ligand exchange reactions. In this method,
203 the limits imposed by the slow exchange kinetics in most ligand exchange reactions
204 are lifted by measuring the free energy difference in the ligand exchange reaction
205 along a user-defined reaction path, taking advantage of the fact that this energy
206 difference is independent of the chosen reaction path. The method implementation is
207 similar to that used by Mei et al. (2013a), and only a short summary is provided here;
208 specific details of employing thermodynamic integration to calculate formation
209 constant of Y(III) chloride complexes are described in section 3.4. We chose to
210 investigate the chloride dissociation reactions, i.e.



212 The reaction paths were defined by constraining the Y(III)-Cl distances (r)
213 step-wise along coordinates that vary from the equilibrium bond (distance r_I) to a
214 distance where the Cl^- ion does not interact with the complex (bulk distance, r_{II}). The
215 mean constraint force ($f(r)$) required to maintain the Y(III)-Cl distance at r was
216 obtained at each step. At each distance r , the simulation was conducted for at least
217 6 ps to sample a significant ensemble of possible configurations of Y(III)-Cl
218 complexes and the surrounding solvent. The Helmholtz free energy of the ligand
219 exchange reaction ($\Delta A_{I \rightarrow II}$) was retrieved by integrating $f(r)$ over the constrained
220 distance (r_I to r_{II}) (Sprik et al., 1998; Sprik & Michiel, 2000):

$$221 \quad \Delta A_{I \rightarrow II} = - \int_{r_I}^{r_{II}} f(r) dr \quad (2)$$

222 The Helmholtz and Gibbs free energies are related by

223
$$\Delta_r G = \Delta A_{I \rightarrow II} + V \int_{P_0}^P dP \quad (3)$$

224 where V is the volume of the box and $\int_{P_0}^P dP$ is the energy contribution due to the
 225 pressure change associated with the ligand exchange at constant volume. As shown in
 226 our previous work (Mei et al. 2013a), the contribution of $\int_{P_0}^P dP$ is small and thus is
 227 neglected. The standard Gibbs free energies of reactions ($\Delta_r G^\ominus$) were calculated from
 228 the Gibbs free energy of reactions ($\Delta_r G$) using

229
$$\Delta_r G^\ominus = \Delta_r G + RT \ln \frac{c_A \gamma_A \cdot c_B \gamma_B}{c_C \gamma_C \cdot c_D \gamma_D} \quad (4)$$

230 where c_i are the concentration of the compounds of reaction $A + B \rightarrow C + D$, and γ_i
 231 are the activity coefficients of each species. By assuming $\gamma_i = 1$ the Gibbs free
 232 energy of reactions of ideal solution ($\Delta_r G^{\ominus,c}$) were obtained. However, in this work,
 233 all simulations were conducted with high relatively concentrations (i.e. 3 m Cl⁻ and
 234 5 m Cl⁻) and high ionic strengths. The activity coefficients of the ions involved in the
 235 ligand exchange reactions were estimated with the B-dot extension of Debye-Hückel
 236 theory (Helgeson, 1981):

237
$$\log \gamma_i = -\frac{z_i^2 A_\gamma I^{1/2}}{1 + a_i B_\gamma I^{1/2}} + \dot{B}_\gamma I \quad (5)$$

238 where z_i is the charge of ion i ; I is the ionic strength in molality; a_i is the ion size
 239 parameter (Å); A_γ and B_γ are defined in Tables 1 and 2 in Helgeson and Kirkham
 240 (1974), \dot{B}_γ is an empirical parameter taken from Oelkers and Helgeson (1990).
 241 Finally, the formation constants (K^\ominus) for the ligand exchange reactions were
 242 calculated from

243
$$\Delta_r G^\ominus = -RT \ln K^\ominus \quad (6)$$

244 where R is the gas constant and T is temperature in Kelvin.

245 The uncertainties of the Gibbs free energies and $\log K^\ominus$ values in *ab initio*
246 thermodynamic integrations were estimated with the method proposed by [Allen and](#)
247 [Tildesley \(1987\)](#). This method has been used by [Rodinger et al. \(2005\)](#) and [Mei et al.](#)
248 [\(2018\)](#) to estimate the uncertainty of thermodynamic integration. The statistical error
249 of each mean constraint force ($\Delta f(r)$) was calculated as the standard deviation of the
250 mean force divided by a statistical inefficiency coefficient; this error includes both the
251 systematic and statistical errors (detail in section 6.4 in [Allen and Tildesley, 1987](#)).
252 The statistical error of $\Delta_r G$ and $\log K^\ominus$ are obtained by accumulating $\Delta f(r)$ over
253 distance ($-\int_{r_I}^{r_{II}} \Delta f(r) dr$). The error of cumulative formation constant was measured
254 using standard propagation rules.

255 2.3 Thermodynamic properties

256 To obtain the formation constants of Y(III)-Cl complexes over a broad
257 temperature and pressure range, we fitted the MD calculation results with the
258 modified Ryzhenko–Bryzgalin (MRB) model ([Ryzhenko et al., 1985](#)):

$$259 \quad \log K_{(T,P)} = -\frac{T_r}{T} \log K_{(T_r,P_r)} + B_{(T,P)} \left(A_{(zz/a)} + \frac{B_{(zz/a)}}{T} \right) \quad (7)$$

260 where K is the formation constant of Y(III)-Cl⁻ complexes at certain T and P ; T_r , P_r
261 are the reference temperature and pressure (*i.e.*, 25 °C and 1 bar), and $A_{(zz/a)}$ and $B_{(zz/a)}$
262 are fitted empirical parameters ([Shvarov and Bastrakov, 1999](#)). The $B_{(T,P)}$ parameter is
263 a species-independent function computed from the dissociation constant of water
264 from [Marshall and Franck \(1981\)](#) (equation (2)). The fitting was done using the

265 OptimC code developed by [Shvarov \(2008, 2015\)](#). The modeling of speciation was
266 conducted using the Matlab code BeerOz ([Brugger, 2007](#)).

267

268 2.4 XAS measurements and data processing

269 Experimental solutions for XAS measurements were prepared by dissolving the
270 appropriate amounts of solids in acidified water. The compositions of the solutions are
271 listed in [Table 3](#). Analytical grade yttrium chloride ($\text{YCl}_3 \cdot 6\text{H}_2\text{O}$), LiCl, NaCl, 37%
272 HCl and MilliQ water were used to prepare the solutions.

273 [Table 3. XAS solution](#)

274 The speciation of yttrium in Cl-rich fluids was characterised via *in-situ* X-ray
275 absorption measurements (XAS) using the high temperature-high pressure cell
276 developed at the Institut Neel/ESRF ([Bruyère et al., 2008](#); [Testemale et al., 2005](#)).
277 This cell consists of an external water-cooled pressure vessel equipped with three
278 beryllium windows enabling collection of simultaneous fluorescence and transmission
279 signals. The sample is contained inside a glassy carbon tube with an internal diameter
280 of 4 mm that is placed inside a small cylindrical resistive heater. Pressure is applied to
281 the sample by two glassy carbon pistons, using helium as a pressure medium, while
282 temperature is recorded by two thermocouples placed within the heating element.
283 Although the thermocouple sits near the sample, it is placed outside the sample and
284 the temperature of the fluid at the beam location needs to be calibrated (e.g., [Brugger](#)
285 [et al., 2007](#); [Etschmann et al., 2010](#); [Liu et al., 2011](#); [Louvel et al., 2017](#)). To do so,

286 the density of pure water is calculated from measurements of the absorbance and
287 X-ray mass attenuation coefficients tabulated by [Chantler \(1995\)](#), and then compared
288 to that from the equation of state of pure water (NIST database, [Lemmon et al., 2000](#)).
289 This method provides a direct measurement of the temperature at the beam position;
290 temperature accuracy is better than 5 °C, and temperature varied within less than 1 °C
291 during the measurements. Pressure was read from a calibrated Bourdin gauge with a
292 precision better than 5 bars; during the experiment, pressure was maintained with
293 1 bar by the PID system ([Bruyère et al., 2008](#)).

294 Yttrium K-edge (17,038 eV) X-ray Absorption Near Edge Structure (XANES)
295 and Extended X-ray Absorption Fine Structure (EXAFS) spectra were collected up to
296 500 °C at 800 bars at the FAME beamline at the European Synchrotron Research
297 Facility (ESRF) in Grenoble, France. The ESRF is a 6.03 GeV ring, and operating in
298 7/8 multi-bunch mode it has a maximum current of 200 mA. FAME is a bending
299 magnet beamline (see [Proux et al., 2005](#) for details), and its double crystal Si(220)
300 monochromator provides an energy resolution of 0.87 eV at the Y K edge energy. A
301 focused beam size of FWHM 220 x 145 μm^2 was used. The incident and transmitted
302 beam intensities I_0 and I_1 were measured with Si diodes, and a Canberra 30 element
303 solid state fluorescence detector was used for detecting fluorescence data. The beam
304 energy was calibrated with an Y foil, such that the maximum of the first derivative
305 was at 17,038 eV.

306 XANES and EXAFS data were analysed with the HORAE package ([Ravel and](#)

307 [Newville, 2005](#)), using FEFF version 9 ([Rehr et al., 2009](#)); E_0 was defined as the
308 maximum of the first derivative. Reported errors in EXAFS parameters ($1-\sigma$) are
309 those calculated by the Artemis program (part of HORAE).

310 *2.5 ab initio XANES simulation*

311 Ab initio XANES simulations were conducted using the FDMNES program
312 ([Joly, 2001](#); [Joly et al., 2009](#)) following the method described in previous studies of
313 metal complexing ([Testemale et al., 2009](#); [Tian et al., 2014](#); [Etschmann et al., 2018](#)).
314 The photo absorption cross sections were calculated using the Finite Difference
315 Method (FDM), which allows accurate calculations especially in the case of
316 low-symmetry structures compared to methods based on the Muffin-Tin
317 approximation, but at a cost of increased computing resources ([Joly, 2001](#); [Brugger, et](#)
318 [al., 2007](#)). To compare with the experimental spectra, the calculations were
319 convoluted with a Lorentzian function of energy-dependent width to reproduce the
320 core-hole lifetime broadening (3.52 eV; [Krause and Oliver, 1979](#)) and the inelastic
321 plasmon interactions with the photoelectron. A Gaussian function (width of 0.87 eV)
322 was employed to account for the energy resolution of the beamline at the convolution.
323 In this study, theoretical spectra were calculated for: 1) crystalline Y_2O_3 solid ([Santos](#)
324 [et al., 2005](#)), as a reference to estimate the accuracy of the simulations; 2) regular
325 predefined square antiprism structures YO_8 ; 3) regular triangular antiprism structures
326 and selected representative Y-O-Cl complexes (with total coordination of 6) from MD
327 calculation at each condition ([Figure 1a,c,d,e](#)); 4) average of at least 50 MD whole

328 boxes at each condition. The radius cutoffs of all calculations were 5 Å.

329 **3. Results**

330 **3.1 Geometrical properties from *ab initio* MD**

331 Unconstrained MD simulations were performed in four solutions with different
332 concentrations of Cl⁻ (Table 2) at 25 °C, 200 °C, 350 °C, 450 °C and 500 °C. The
333 simulation results are summarized in Table 4. The snapshots of different Y(III)-Cl⁻
334 complexes and pair distribution functions of Y-O and Y-Cl of each solutions are
335 shown in Figure 1 and 2.

336 Table 4. simulation results from *ab initio* MD

337 Figure 1. snapshots of Y(III)-Cl

338 Figure 2. RDF

339 In box 1 (Y:Cl = 1:3), at ambient conditions (25 °C, 1 bar), two jobs were
340 conducted to test the stability of Y(III)-Cl complexes. In job 1a(1), the simulation
341 started with [Y(OH₂)₈]³⁺, and this configuration remained stable during the whole
342 simulation. In job 1a(2), the initial configuration was [Y(OH₂)₇Cl]²⁺ and Cl⁻
343 dissociated from Y(III) in less than 1 ps (Fig. 3a), and [Y(OH₂)₈]³⁺ remained stable
344 for the rest of the simulation. The eight water molecules in the first shell of Y³⁺
345 formed an antiprism structure (Fig. 1a, 1b), which is consistent with previous MD
346 and XANES studies (Table 1). The Y-O bond distance (2.42 Å) from MD is also
347 within the range of values from literature (2.36 Å to 2.51 Å, reference in Table 1),
348 and close to our XAS results (2.37 to 2.39 Å, section 3.2). No evidence of Cl⁻

349 bonding with Y(III) was found for this Cl concentration. The oscillations of Y-O
350 bond distance indicate Debye-Waller factors of 0.0106 \AA^2 and 0.0107 \AA^2 for
351 job 1a(1) and 1a(2), respectively.

352 Simulation 1b was run at 200 °C and 800 bar for a total of 30.17 ps, starting with
353 the $[\text{Y}(\text{OH}_2)_6\text{Cl}_2]^+$ complex as the initial configuration. The calculated hydration
354 number is 5.2. One Cl^- dissociated from Y(III) from 14.6 ps to 22.6 ps (Fig. 3b),
355 which leads to an average chloride coordination number of 1.8 over the simulation
356 time. The RDFs in Fig. 2 indicate Y-Cl and Y-O bond distances of 2.65 Å and 2.36 Å,
357 respectively. At 350 °C, 800 bar (simulation 1c), the total coordination number of
358 Y(III) further decrease to 6.5. Two water molecules bonded with Y(III) in the initial
359 configuration detached from Y(III) within the first picosecond. The two Cl^- remained
360 bonded to Y(III) at an average distance of 2.62 Å over the 23.87 ps simulation, *i.e.* no
361 ligand change was observed. At 500 °C, 800 bar and 1000 bar (simulation 1d and 1e),
362 the final species was $[\text{Y}(\text{OH}_2)_5\text{Cl}_3]^0$. The Y-Cl and Y-O bond lengths (Fig. 2) were
363 2.55 – 2.56 Å and 2.44 Å, respectively.

364 **Figure 3. Y-Cl distance over time**

365 Simulations of box 2, 3, 4 were conducted with different Cl^- concentrations
366 (Y:Cl = 1:4, 1:5, 1:2.5), and run at the same P-T conditions as box 1 except room
367 temperature. The bond distances and total coordination numbers calculated within
368 each box are consistent at each P-T condition (Table 2). At 200 °C, the coordination of
369 Cl^- in box 4a is 1, whereas it is ~2 in the other boxes, suggesting an effect of

370 increasing Cl contents on the stability of Y-Cl complexes. At 500 °C, 1000 bar,
371 coordination of Cl⁻ is 2 for simulation 2c. In simulation 4c (Cl⁻ concentration of 5 m),
372 at 450 °C, 800 bar, the initial complex [Y(OH₂)₅Cl₃]⁰ lost two waters within the first
373 1.5 ps, and became [Y(OH₂)₃Cl₃]⁰, which remained stable for the last 15.41 ps.
374 Interestingly, the coordination number of Cl⁻ reaches 4 in simulation 3c (Y:Cl = 1:5;
375 500 °C, 1000 bar), with the average number of water molecules down to 1.7.

376 In general, the total coordination number decreases from 8 to 6 and Y(III)-Cl⁻
377 complexes become more stable as temperature increases from 25 °C to 500 °C. At
378 high temperature (500 °C), more chlorides are coordinated with Y(III) and hydration
379 number decreases with more Cl⁻ complexes for a given P-T condition, different
380 complexes were observed from MD simulations in each box. This reflects the
381 different Y:Cl ratios in the boxes, but also the fact that the simulations in each box at
382 the same P-T condition were started from different initial configurations, and due to
383 the short time range of MD relative to ligand exchange rates, the final species
384 detected from MD could be meta-stable.

385 3.2 XAS data refinement

386 3.2.1 XANES

387 XANES spectra are mainly a function of the oxidation state of the target atom,
388 and the atomic number (e.g. O versus Cl) and geometrical arrangement of the ligands
389 coordinated to the target atom. The edge position (red-shifted as the formal charge on
390 the target atom decreases) and intensity of the white line (most intense band at the

391 edge position) can change as the ligands vary depending on composition and
392 temperature (Bunker, 2010, Penner-Hahn, 2005; Etschmann, et al., 2017).

393 Yttrium XANES spectra show the classic “dehydration” effect with increasing
394 temperature and salinity (e.g. Brugger et al., 2016). This is reflected by a decrease of
395 the white line intensity and a shift to lower energy of the edge position, as shown in
396 Figure 4 where the spectra obtained on Sol3 (0.01 m YCl_3) and Sol5 (9.5 m LiCl +
397 0.5 m HCl) at 25 and 400 °C are compared. The effect of increasing temperature and
398 salinity combine so that the decrease of the white line intensity and energy shift are
399 more pronounced for Cl-rich Sol5. Note that that the small energy shift is more
400 clearly observed in the first derivative of the XANES spectra (Fig 4(b)).

401 Figure 4. XANES on highest Cl solutions

402 The qualitative discussion above was supported using *ab initio* XANES
403 simulations. As shown in Figure 6(a), the positions of the spectral features are well
404 reproduced by the calculations for the $Y_2O_3(s)$ standard, except for the band at
405 ~65.9 eV, which is shifted by ~8 eV. The relative intensities are well reproduced, but
406 the total intensity of the white line is lower in the simulations than in the experiments.
407 Figure 6(b) explores the shifts of the spectral features resulting from changes in bond
408 distance and angles in a high symmetry $[YO_8]$ moiety. The angle affects the shape of
409 the white line (intensity ratio of the two main bands contributing to the white line),
410 and the bond distance causes a shift in the position of the white line. The MD
411 structures -and most likely the real species in solution- reflect dynamic distortions of

412 the antiprism structure, which results in the small but significant differences in
413 calculated XANES spectra (Figure 6(c)). In order to account for this, we calculated
414 the averages from ≥ 50 frames over the simulation times for a number of simulations
415 (Figure 6(d)). In general, the results of FDMNES of MD boxes and experiments agree.
416 Importantly, the effects of temperature and salinity on the spectra are well reproduced
417 by the simulations. It is noticeable that the Y-O distance at room temperature from
418 MD and XAS is 2.42 Å and 2.37 Å, respectively; therefore, minor discrepancies
419 between the spectra from FDMNES of MD boxes and spectra from experiments are
420 reasonable.

421 Figure 6. XANES from FDMNES.

422 3.2.2 EXAFS

423 EXAFS spectra represent a final state interference effect arising from the
424 scattering of the outgoing photoelectron from neighboring atoms, and are therefore
425 mainly sensitive to the nature (atomic number), quantity and distance of the
426 neighboring atoms.

427 The amplitude reduction factor (S_0^2) accounts for processes that contribute to
428 the X-ray absorption coefficient but not to the EXAFS (Roy et al. 1997), and is
429 determined by fitting a compound with similar molecular structure as the unknown. In
430 our case, S_0^2 was determined by fixing the coordination of a low salinity solution,
431 Sol6 (0.75 m HCl) at 30 °C to eight oxygen atoms, because the structure of the Y(III)
432 aqua ion has been well characterized by previous studies (Ikeda et al., 2005a; Ohta et

433 al., 2008; Liu, X. et al., 2012; Lindqvist-Reis et al., 2000). A S_0^2 value of 0.73 was
434 determined by refining the S_0^2 , bond lengths and EXAFS Debye-Waller factors for
435 this solution; this value is consistent with theoretical constraints (Roy et al. 1997) and
436 was used in all subsequent refinements.

437 The YOCl(s) structure (Zachariasen, 1949, ICSD # 31667) was used to generate
438 the O and Cl paths. For each solution, all spectra were fitted simultaneously, resulting
439 in one ΔE_0 and one set of goodness-of-fit parameters for each solution (Table 5). The
440 total ligand number (one oxygen path and one chlorine path) was restrained to be
441 between 6 and 8; attempts to decrease the minimum coordination number to 5 resulted
442 in statistically inferior fits. The Debye-Waller σ^2 was fitted at 25 °C for each solution,
443 with the constraint that it should then double from 30 to 500 °C, as determined from
444 MD simulations. Note that the Debye-Waller values determined from the *ab initio*
445 MD simulations are higher than the experimental ones, as a result of the short
446 sampling times in *ab initio* MD, that tend to overestimate structural disorder. However,
447 the relative changes in fitted Debye-Waller values are likely to be accurate, especially
448 since the vibrational characteristics of the complexes are well reproduced by *ab initio*
449 MD (e.g. Mei et al. 2013b and references therein). In practice this was applied by
450 increasing the EXAFS Debye-Waller by a factor of 1.2 for each ~100 °C increase in
451 temperature. The EXAFS Debye-Waller for Cl⁻ was defined to be the same as that for
452 O, in part to reduce the number of variables and in part due to the (usually) low Cl:O
453 ratio.

454 The Y k-space and Fourier transformed (R-space) EXAFS data are shown in
455 [Figure 5](#) and the fit results are tabulated in [Table 5](#). As a general rule, the data show a
456 small shift in k-space with increasing temperature. In R-space, the magnitude of the
457 peak in the radial distribution function decreases, and the peak position shifts to a
458 longer distance with increasing temperature, consistent with decreasing total
459 coordination numbers and increasing Cl:O ratio. This evolution is also seen in the
460 fitted structural parameters ([Table 5](#)): the largest change is recorded in Sol5 (9.5 m
461 LiCl+0.5 m HCl), where the total ligand number decreased from 7.6 (30 °C) to 6.0
462 (450 °C), with the Cl:O ratio increasing from 0.0 to 1.0. Attempts were made to
463 increase the Cl:O ratio to 2, i.e. $4xCl+2xO$, in-line with the results from MD, however
464 this skewed the fit peak significantly, resulting in a poor overall fit to the data.

465 [Figure 5. EXAFS](#)

466 [Table 5 EXAFS result](#)

467 In contrast, the total coordination number does not decrease as much in low
468 salinity solutions, e.g. from 8 to 7 (Sol3, 30 to 400 °C) and 8 to 7.7 (Sol6, 30 to
469 400 °C), while the Cl:O ratio increases from 0 to ~0.6 and ~0.4 (Sol6 and Sol3
470 respectively).

471 3.3 Estimating formation constants by *ab initio* thermodynamic 472 integration

473 The MD simulations described above provided qualitative information about the
474 nature and geometry of the Y-Cl complexes in chloride solutions to 500 °C. However,

475 thermodynamic properties of Y-Cl complexes are necessary to determine their relative
476 stabilities and the mobility of Y(III) under hydrothermal conditions.

477 **Figure 7. FES of YCl^{2+}**

478 **Figure 7** shows an example of employing *ab initio* MD to calculate the free
479 energy surface of the reaction:



481 The Helmholtz free energy $\Delta A_{I \rightarrow II}$ of reaction (8) was calculated by integrating the
482 constraint mean force ($f(r)$) between Y^{3+} and Cl^{-} (**equation (1)**) from the equilibrium
483 position (2.65 Å), which corresponds to the equilibrated Y(III)- Cl^{-} bond distance in
484 the previous unconstrained MD simulations, to a non-interacting position (6.00 Å). An
485 external force is required to eliminate the interactions between Y^{3+} and Cl^{-} to maintain
486 the targeted Y(III)- Cl^{-} distance. When the Y(III)- Cl^{-} distance was between 3.1 Å and
487 3.55 Å, the absolute value of $f(r)$ varied from zero to a minimum of -45.14 kJ/mol•Å
488 to near zero (3.32 kJ/mol•Å), and the free energy reached its maximum (23.50 kJ/mol),
489 which means at this stage, one water molecule associated with yttrium, i.e. reaction (7)
490 results in the replacement of one chloride ion by one water molecule. The mean
491 constraint forces become positive at distances between 3.4 Å and 5.4 Å,
492 which reflects the activation barrier of the ion exchange reaction. It can be seen from
493 **Figure 7** that when the Y(III)- Cl^{-} distance is beyond 4.5 Å, the constraint mean force
494 is very small (9.01 kJ/mol•Å, 2.03 kJ/mol•Å, -2.69 kJ/mol•Å, and -9.45 kJ/mol•Å at
495 distances of 4.5 Å, 4.9 Å, 5.4 Å, and 6 Å, respectively), as expected as chloride-Y(III)

496 interaction become vanishingly small. The integration of the Y(III)-Cl⁻ mean
497 constraint force over the reaction path gives a free energy for reaction (7) of $3.92 \pm$
498 8.85 kJ/mol.

499 [Figure 8. free energy surface of all calculation](#)

500 **Figure 8** shows the free energy surface of the stepwise dissociation of
501 Y(III)-chloride complexes in solutions with different Cl⁻ concentrations (box 1 and 3)
502 at 200 °C, 800 bar; 350 °C, 800 bar; and 500 °C, 1000 bar. In order to achieve good
503 sampling of the constraint force, one more free chloride was set in the system at a far
504 distance (6 Å) to conduct thermodynamic integration, i.e. in box 1, which contains
505 three Cl⁻, the free energies of formation of YCl_n³⁻ⁿ complexes were calculated only
506 when n=1 and 2; in box 3, which contains five Cl⁻, $\Delta_r G$ of YCl_n³⁻ⁿ (n = 1,2,3,4) were
507 measured (**Table 6**).

508 [Table 6. \$\Delta G\$ and \$\log K^\ominus\$ of two boxes](#)

509 Comparing results from box 1 and box 3, the calculated $\Delta_r G$ for the association
510 reactions of YCl²⁺ and YCl₂⁺ vary because of the difference of Cl⁻ activity in each box.
511 After activity correction, the **value of** standard Gibbs free energies ($\Delta_r G^\ominus$) of each
512 ligand exchange reaction were close (four are ≤ 2.2 kJ/mol, and two ≤ 3.8 kJ/mol) at
513 the same P-T condition. The stepwise formation constants, $\log K^\ominus$, that were obtained
514 from the two boxes at each P-T condition, their errors and averages are presented in
515 **Figure 9**. The consistency of the two $\log K^\ominus$ values from the two boxes validates the
516 thermodynamic integration method for calculating the energetics of these ligand

517 exchange reactions, demonstrating that the values do not depend on the composition
518 or size of the simulation box, and that the sampling time of the simulations is
519 sufficient. In this study, we take the average stepwise formation constant of the two
520 boxes as the final stepwise formation constant of the YCl^{2+} and YCl_2^+ association
521 reactions at 200 °C, 800 bar; 350 °C, 800 bar; and 500 °C, 1000 bar. The error of the
522 average formation constant was measured using the standard propagation rules. The
523 stepwise ($\log K^\ominus$) and cumulative ($\log \beta^\ominus$) formation constants of Y(III)-chloride
524 complexes are summarized in [Tables 6 and 7, respectively](#).

525 [Figure 9](#). average $\log K^\ominus$ and each $\log K^\ominus$ within error bar in different boxes with T.
526 [Table 7](#). $\log \beta$ of Y(III)-Cl.

527 The calculated $\log \beta$ in [Table 7](#) are employed to fit the modified MRB model
528 ([eq. 6](#)) to extrapolate the formation constant of Y(III)-Cl⁻ associate reactions to higher
529 temperature and pressure. We fit two parameters in [equation \(6\)](#) for each species to
530 minimize the uncertainties. The $\log \beta_{(25^\circ C, 1 \text{ bar})}^\ominus$ of YCl^{2+} was chosen from Luo &
531 Byrne ([2001](#)), which is the only available experiment data of Y(III)-Cl complexes,
532 two parameters (*A* and *B*) were fitted; but for YCl_2^+ , YCl_3 , and YCl_4^- , the $B_{(zz/a)}$ are set
533 to zero, and $\log \beta_{(25^\circ C, 1 \text{ bar})}^\ominus$ and the *A* parameter were fitted. The fitted MRB
534 parameters are listed in [Table 8](#). The extrapolated logarithm of formation constants of
535 Y(III)-Cl complexes at 25 °C to 500 °C, saturated pressure (P_{sat}) to 5000 bar from
536 MRB, are listed in [Table 9](#).

537 [Table 8](#). MRB parameters

538 Table 9. $\log\beta$ from MRB

539 **4 Discussion**

540 4.1 Stability of Y(III)-Cl complexes

541 In general, the speciation derived from the unconstrained MD (section 3.1)
542 agrees well with XAS fits (section 3.2) (Figures 10 and 11). The XANES spectra
543 calculated from *first principles* from the MD simulations reproduce the experimental
544 spectral features and their changes with temperature and salinity well (Figure 6d). The
545 experiments and theoretical approaches both reveal the same trends of Y(III)-Cl
546 complexes along with temperature, with more Cl⁻ bonding with Y(III) at high
547 temperatures.

548 Figure 10. Coordination of O and Cl from XAS and MD

549 The formation constants obtained from thermodynamic integration enable us to
550 predict Y(III)-Cl⁻ speciation at different PT and Cl⁻ concentrations and provide the
551 first self-consistent speciation model for Y(III) chloride complexing from
552 room-temperature to high-temperature hydrothermal conditions. The results indicate
553 that the formation of Y(III)-chloride complexes can affect the mobility of yttrium
554 under hydrothermal conditions and also the distribution of REE elements in REE
555 deposits.

556 Figure 11 shows the distribution of Y(III) chloride complexes at different T, P
557 and pH, as a function of total chloride in the solution. The number of coordinated
558 chloride ligands from XAS Sol3 (0.03 m Cl⁻), Sol5 (2.5 m Cl⁻) and Sol7 (5 m Cl⁻) are

559 also plotted in **Figures 11a,c** and **11d,e,f** for comparison (right axis, orange dots). At
560 200 °C and $\text{pH}_{200^\circ\text{C}} = 0.3$ and 3, the hydrated Y^{3+} ion remains the dominant form of
561 Y(III) even in high salinity solutions (**Figure 11a,b**). At 350 °C, the stability of the
562 different species not only varies with Cl^- concentrations, but also with pH, so that the
563 predominant species remains $\text{YCl}_3(\text{aq})$ from $n(\text{Cl}^-) > 0.4 \text{ m}$ at $\text{pH}_{350^\circ\text{C}} = 0.3$, but
564 changes to YCl_4^- at $\text{pH}_{350^\circ\text{C}} = 3$. At 500 °C, 1000 bar (**Figure 11e,f**), the dominant
565 species is YCl_4^- for $\text{pH}_{350^\circ\text{C}} = 3$, and $\text{YCl}_3(\text{aq})$ for $\text{pH}_{350^\circ\text{C}} = 0.3$. The coordination
566 number of Cl^- is also 3 in XAS Sol 5 at 450 °C, whose Cl^- concentration is 10 m but
567 also with high acidity. This decrease of Cl neighbors is likely due to the association of
568 HCl at high temperature (**Mei et al., 2018; Tagirov et al., 1997**) that leaves less free
569 Cl^- available to bond with Y(III) in very acidic solutions.

570 While the general trends defined by (i) XAS, (ii) unconstrained MD (**Figure 10**)
571 and (iii) thermodynamic modelling (**Figure 11**) derived from thermodynamic
572 integration are consistent with each other, small discrepancies in the coordination
573 number between the methods are noticeable. For example, at 200 °C, both
574 unconstrained MD and the thermodynamic model indicate that there are Cl^-
575 bonding with Y(III) when $n(\text{Cl}^-) > 2.5 \text{ m}$ (1 to 2 in MD and ~ 0.5 in the
576 thermodynamic model), whereas XAS analysis **reveals very minor chlorine**
577 **complexation in Sol 7 and no chlorine complexation in Sol 6 (Table 5 Sol6)** (i.e.,
578 below detection limit of $\sim 0.3 \text{ Cl}^-$). At 350 °C, the thermodynamic model suggests
579 higher order Y(III)-Cl species than MD and XAS. This is because:

580 (1) thermodynamic integration with the MD method tends to overestimate the
581 formation constants of complexes, and especially higher order complexes, at
582 relative low temperature due to the low kinetic energy of the ions and water
583 molecules, resulting in poor sampling of the force (Mei, et al., 2018). This
584 translates into larger errors on the $\log\beta$ values at 25 °C than at 300 °C, for example
585 (Mei et al., 2013a, 2015a). Therefore, the $\log\beta$ of Y(III)-Cl species at 200 and
586 350 °C from MD that was employed to model Y(III)-Cl distribution in this study
587 could still be higher than actual, and consequently leads to the overestimation of the
588 coordination of Y(III)-Cl. (2) As discussed above, due to the limits of CPU
589 resources, the simulation time of unconstrained MD is not sufficient to observe the
590 stable species, so species reported in Figure 10 could be meta-stable. (3) The
591 experimental solutions are more acidic than those in the simulations, and HCl
592 association could explain the lower Y-Cl bonding compared to other solutions and
593 to the MD simulations (Fig. 11).

594 Overall, Y(III) mobility in hydrothermal conditions is expected to increase
595 because of the increased stability of higher-order Y(III)-Cl with increasing
596 temperature and Cl⁻ concentration.

597 Figure 11. Y-Cl species with n(Cl⁻) at different T and pH

598 4.2 REE-Cl⁻ complexation in hydrothermal fluids

599 The Y³⁺ ionic radius is close to that of Ho³⁺, and Y has consequently frequently
600 been grouped with heavy REE. However, differences in their aqueous speciation has

601 the potential to lead to different mobility of the REE and Y in natural hydrothermal
602 systems. Up to now, this possibility could not be investigated due to the lack of
603 thermodynamic properties for Y(III)-Cl complexes (Migdisov et al., 2016). The
604 formation constants of YCl^{2+} and YCl_2^+ at 150, 200 and 250 °C, and P_{sat} have been
605 derived from the fitted MRB parameters of Y(III)-Cl complexes (Table 9) and can
606 now be compared to other REE data (Table 1, Migdisov et al., 2009) (Figure 12). At
607 150 °C, the formation constant for YCl^{2+} is close to zero, much smaller than those
608 reported by Migdisov et al. (2016) for $REECl^{2+}$, even when considering the heaviest
609 REE (Lu, which is 1.14). At 200 °C, yttrium behaves similar to the heavy REE
610 (HREE) element (Lu) in Cl-rich solutions. In this condition, the LREE are more stable
611 in solution and mobile than HREE, including Y(III). Nevertheless, the formation
612 constants of Y-Cl complexes increase faster than those of other HREE with increasing
613 temperature, so that at 250 °C Y(III)-Cl complexes are more stable than HREE, and Y
614 behaves more like the LREE (La).

615 *Figure 12. Compare of $\log\beta$ of REE & Y values*

616 Differences between Y and the HREE thermodynamics under hydrothermal
617 conditions provide a likely explanation for the fractionation of Y from Ho reported in
618 some hydrothermal REE minerals (Bau and Dulski, 1995). The distribution of Ho(III)
619 and Y(III) species as a function of Cl^- concentration at 250 °C, P_{sat} are presented in
620 Figure 13, with the data presented in Table 9 for Y(III) and measured by Migdisov et
621 al (2009) for Ho^{3+} , $HoCl_2^+$ and $HoCl^{2+}$, and predicted from Haas et al. (1995) for

622 $\text{HoCl}_3(\text{aq})$. As thermodynamic data for Ho(III) are only available until trichloro
623 complexes, YCl_4^- is not considered in **Figure 13**. At 250 °C, the dominant species for
624 Y(III) in Cl-bearing solution is YCl_2^+ , whereas for Ho(III), it remains HoCl^{2+} until Cl
625 concentration reaches ~4.5 m and becomes HoCl_2^+ afterwards. Such slight differences
626 could ultimately account for contrasting REE(Y) distribution in hydrothermal REE
627 deposits. However, the minerals deposited from hydrothermal fluids contain many
628 components, i.e. F^- , SO_4^{2-} , CO_3^{2-} . The behaviors of REE with most of these ligands
629 are unclear but can affect the process of precipitation; therefore, further studies are
630 required to quantitatively model how REE minerals deposits in hydrothermal process.

631 **Figure 13. Y-Cl/Ho-Cl species distribution with Cl⁻ concentration at 250 °C, P_{sat}**

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643 **Research data**

644 Relevant research data are in attached spreadsheet.

645

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- 926

Table 1 Structure of aqueous Y(III) at room temperature (~25 °C) and hydrothermal conditions in literature.

	Methods	conditions	hydration	Y-O distance (Å)	reference	note
Experimental studies	XRD	1.094 m Y(ClO ₄) ₃	8.0 ± 0.3	2.365	Johansson and Wakita, 1985	
	AXS	0.5 M YCl ₃	8.1 ± 0.3	2.51 ± 0.02	Matsubara et al., 1990	
		1 M YCl ₃	8.2 ± 0.5	2.46 ± 0.02		
	EXAFS	0.5 M YCl ₃	7.5	2.33	Marques et al., 1992	No Y-Cl bonds were found in either Raman and EXAFS
		1.0 M YCl ₃	8.5	2.33		
		1.2 M YCl ₃	7.5	2.34		
	EXAFS	(a) 0.1 M YCl ₃	9.1	2.36	Ragnarsdottir et al., 1998	No Y-Cl bonding and hydration decreases by 0.6, 2.6 and 1.8 with temperature increasing up to 340 °C for (a) and 250 °C for (b) and (c)
		(b) 0.1 M YCl ₃ + 0.23M HCl + 1M NaCl	10.2	2.36		
		(c) 0.1 M YCl ₃ + 0.23M HCl + 2M NaCl;	10.3	2.37		
		EXAFS&LXAS	1.71 M Y(ClO ₄) ₃ in HClO ₄	8	2.368 ± 0.003	Lindqvist-Reis et al., 2000
	EXAFS	0.005 ~ 2 m YBr ₃	8	2.353 ± 0.001	Diaz-Moreno et al., 2000	
	EXAFS	0.1 M YCl ₃ + 0.25 M HCl/14 M LiCl	9.7/9.6		Allen et al., 2000	
	EXAFS & EPSR	1 m YCl ₃	7.5 ± 0.5	2.33	Bowron and Diaz-Moreno, 2007	0.8 ± 0.2 Y-Cl
	RAMAN/DFT	3.2 M YCl ₃ and 2.03 M YCl ₃ + (0, 1, 4M HCl)	8-n	2.366	Rudolph and Irmer, 2015	Square antiprism, with <u>n</u> Y-Cl bonds (n = 1,2); n=0 when Cl ⁻ concentration < 0.2 m
Theoretical studies	DFT	1 m YCl ₃	8	2.38	Ikeda et al., 2005a, b	Square antiprism structure
	AIMD	1 Y ³⁺ in 63 waters' box	8		Liu et al., 2012	

Table 2. Simulation details of *ab initio* MD.

Box No.	Solution composition	Job No.	T (°C)	P (bar)	Box size (Å ³)	Density (g/cm ³)
1	55 H ₂ O, Y ³⁺ , 3Cl ⁻ (3 m Cl ⁻)	1a(1,2)	25	1	12.048	1.127
		1b	200	800	12.366	1.042
		1c	350	800	12.999	0.897
		1d	500	800	14.228	0.684
		1e	500	1000	13.940	0.727
2	55 H ₂ O, Y ³⁺ , 4 Cl ⁻ , 1 Na ⁺ (4 m Cl ⁻)	2a	200	800	12.443	1.073
		2b	350	800	13.032	0.934
		2c	500	1000	14.122	0.734
3	55 H ₂ O, Y ³⁺ , 5 Cl ⁻ , 2 Na ⁺ (5 m Cl ⁻)	3a	200	800	12.526	1.101
		3b	350	800	13.085	0.966
		3c	500	1000	13.865	0.812
4	111 H ₂ O, Y ³⁺ , 5 Cl ⁻ , 1 Na ⁺ , H ⁺ (2.5 m Cl ⁻)	4a	200	800	15.577	1.006
		4b	350	800	16.412	0.860
		4c	450	800	17.368	0.726

Table 3. XAS solution compositions.

Sample composition	Serial Number	condition
0.01 m YCl_3 + 0.75 m HCl	Sol 6	30 °C – 400 °C, 800 bar
0.01 m YCl_3 + 10 m LiCl + 0.5 m HCl	Sol 5	30 °C – 450 °C, 800 bar
0.01 m YCl_3 + 2 m NaCl + 0.5 m HCl	Sol 7	30 °C – 350 °C, 800 bar
0.01 m YCl_3 + 1 drop HCl	Sol 3	30 °C – 400 °C, 800 bar
1 m YCl_3	Sol 12	30 °C – 450 °C, 800 bar

Table 4. Simulation results by *ab initio* MD, error of coordination was measured from radial distribution function at distance range of 2.75 Å - 3.25 Å for Y-O and 2.9 Å – 3.2 Å for Y-Cl; Debye-Waller can only be measured when there is no ligand exchange in the simulation.

No	T (°C), P (bar)	Time (ps)	Initial configuration	Stable species	Y-Cl			Y-O			CN _{tot}
					N	r(Å)	σ ² (Å ²)	N	r(Å)	σ ² (Å ²)	
1a(1)	25, 1	25.37	[Y(H ₂ O) ₈] ³⁺	[Y(OH ₂) ₈] ³⁺				8	2.42	0.0106	8
1a(2)	25, 1	23.02	[Y(H ₂ O) ₇ Cl] ²⁺	[Y(OH ₂) ₈] ³⁺				8	2.42	0.0107	8
1b	200, 800	30.17	[Y(H ₂ O) ₆ Cl ₂] ⁺	[Y(H ₂ O) _{5.5} Cl _{1.8}] ^{1.2+}	1.7(0.1)	2.65		5.2(0.1)	2.36		7
1c	350, 800	23.87	[Y(H ₂ O) ₇ Cl ₂] ⁺	[Y(H ₂ O) _{4.5} Cl ₂] ⁺	2(0.1)	2.62	0.0172	4.5(0.1)	2.36		6.5
1d	500, 800	22.43	[Y(H ₂ O) ₅ Cl ₃] ⁰	[Y(OH ₂) ₃ Cl ₃] ⁰	3(0.2)	2.55	0.0155	3(0.2)	2.44		6
1e	500, 1000	26.94	[Y(H ₂ O) ₅ Cl ₃] ⁰	[Y(OH ₂) ₃ Cl ₃] ⁰	3	2.56	0.0200	3	2.44		6
2a	200, 800	20.30	[Y(H ₂ O) ₇ Cl ₂] ²⁺	[Y(H ₂ O) _{5.3} Cl _{1.8}] ^{1.2+}	1.8(0.1)	2.6		5.2(0.1)	2.35		7.1
2b	350, 800	19.21	[Y(H ₂ O) ₇ Cl ₂] ⁺	[Y(H ₂ O) ₅ Cl ₂] ⁺	1.9(0.1)	2.66	0.0236	4.9(0.2)	2.38		7
2c	500, 1000	16.23	[Y(H ₂ O) ₆ Cl ₂] ⁺	[Y(H ₂ O) _{4.2} Cl ₂] ⁺	2(0.1)	2.56		4.1(0.2)	2.33		6.2
3a	200, 800	16.06	[Y(H ₂ O) ₇ Cl] ²⁺	[Y(OH ₂) ₅ Cl ₂] ⁺	2(0.1)	2.67	0.0168	5(0.1)	2.38		7
3b	350, 800	17.52	[Y(OH ₂) ₄ Cl ₃] ⁰	[Y(OH ₂) _{3.9} Cl _{2.3}] ²⁺	2.3(0.1)	2.61		3.9(0.1)	2.35		6.2
3c	500, 1000	17.01	[Y(H ₂ O) ₃ Cl ₃] ⁰	[Y(OH ₂) ₂ Cl ₄] ⁻	4(0.1)	2.59	0.0194	1.7(0.3)	2.45		5.7
4a	200, 800	22.35	[Y(H ₂ O) ₈ Cl] ²⁺	[Y(H ₂ O) ₆ Cl] ²⁺	1	2.63	0.0116	6(0.1)	2.34	0.0155	7
4b	350, 800	16.92	[Y(OH ₂) ₅ Cl ₃] ⁰	[Y(OH ₂) _{3.7} Cl ₂] ⁺	2	2.63	0.0165	4.1(0.1)	2.36		6.1
4c	450, 800	16.91	[Y(OH ₂) ₅ Cl ₃] ⁰	[Y(OH ₂) ₃ Cl ₃] ⁰	3	2.56	0.0191	3.1(0.2)	2.37		6.1

Table 5. EXAFS results.

T (°C)	nO	rO	nCl	rCl	n _{tot} (first shell)	nY	rY	ss	k-range	R-range
Sol 6			E ₀ = 2.2(4)	$\chi^2_{\text{red}} = 661$	r-factor = 0.045					
30	8.0 (fix)*	2.369(7)			8	1 (fix)	4.78 (fix)	0.0041(8)	2<k<11	1.1<R<4.5
200	7.0(4)	2.391(8)	(1)		7	1 (fix)	4.78 (fix)	1.4*ssO1	2<k<11	1.1<R<4.5
300	6.4(1)	2.39(1)	0.6(5)	2.60 (fix)	7	1 (fix)	4.78 (fix)	1.6*ssO1	2<k<11	1.2<R<4.5
350	5.2(4)	2.36(1)	1.8(4)	2.68(2)	7	1 (fix)	3.94 (fix)	1.7*ssO1	2<k<10	1.2<R<4.5
400	4.4(4)	2.34(1)	2.6(4)	2.64(1)	7	1 (fix)	3.94 (fix)	1.8*ssO1	2<k<10	1.2<R<4.5
Sol 5			E ₀ = 3.3(4)	$\chi^2_{\text{red}} = 523$	r-factor = 0.037					
30	7.6(4)	2.388(7)			7.6	1 (fix)	4.74 (fix)	0.0023(3)	2<k<11	1.15<R<4.5
200	6.2(4)	2.398(8)	1.0 (fix)	2.66 (fix)	7.2	1 (fix)	4.74 (fix)	1.4*ssO1	2<k<9	1.15<R<4.5
300	4.0(2)	2.356(8)	2.0(1)	2.68(1)	6	1 (fix)	3.99 (fix)	1.6*ssO1	2<k<10	1.15<R<4.5
350	3.4(1)	2.331(1)	2.6(1)	2.652(9)	6	1 (fix)	3.99 (fix)	1.7*ssO1	2<k<10	1.15<R<4.5
400	3.0(3)	2.338(1)	3.0(1)	2.657(8)	6	1 (fix)	3.99 (fix)	1.8*ssO1	2<k<10	1.15<R<4.5
450	3.0 (fix)	2.326(1)	3.0 (fix)**	2.649(8)	6	1 (fix)	3.99 (fix)	1.9*ssO1	2<k<10	1.15<R<4.5
Sol 7			E ₀ = 2.2(4)	$\chi^2_{\text{red}} = 632$	r-factor = 0.043					
30	8.0(5)	2.362(8)			8	1 (fix)	4.78 (fix)	0.0041(6)	2<k<11	1.0<R<4.5
200	7.3(4)	2.389(8)	(1)		7.3	1 (fix)	4.78 (fix)	1.4*ssO1	2<k<11	1.0<R<4.5
300	5.8(4)	2.38(1)	1.5(4)	2.70(2)	7.3	1 (fix)	3.99 (fix)	1.6*ssO1	2<k<11	1.0<R<4.5
350	5.1(3)	2.36(1)	2.1(3)	2.67(1)	7.2	1 (fix)	3.99 (fix)	1.7*ssO1	2<k<11	1.0<R<4.5
Sol 3			E ₀ = 1.8(4)	$\chi^2_{\text{red}} = 673$	r-factor = 0.039					
30	8.0(1)	2.378(8)			8	1 (fix)	4.74 (fix)	0.0050(4)	2<k<11.5	1<R<4.5
200	7.8(3)	2.384(7)	(1)		7.8	1 (fix)	4.74 (fix)	1.4*ssO1	2<k<11	1<R<4.5
300	6.9(3)	2.38(1)	0.9(3)	2.69(6)	7.8	1 (fix)	3.99 (fix)	1.6*ssO1	2<k<11	1<R<4.5
350	6.4(4)	2.37(1)	1.3(4)	2.68(3)	7.7	1 (fix)	3.99 (fix)	1.7*ssO1	2<k<10	1<R<4.5
400	5.7(6)	2.34(1)	2.0(6)	2.64(3)	7.7	1 (fix)	3.99 (fix)	1.8*ssO1	2<k<9	1<R<4.5
Sol 12			E ₀ = 0.3(5)	$\chi^2_{\text{red}} = 7725$	r-factor = 0.031					
30	8.0(1)	2.374(6)			8	1 (fix)	4.74 (fix)	0.0042(4)	2<k<12	1.1<R<4.0
200	7.9(5)	2.364(7)	0.6(4)	2.60 (fix)	8.5	1 (fix)	4.74 (fix)	1.4*ssO1	2<k<12	1.1<R<4.0
300	6.2(4)	2.36(1)	1.6(4)	2.69(2)	7.8	1 (fix)	3.99 (fix)	1.6*ssO1	2<k<10	1.1<R<4.0

350	5.7(5)	2.36(1)	2.1(4)	2.67(2)	7.8	1 (fix)	3.99 (fix)	1.7*ssO1	2<k<10	1.1<R<4.0
400	5.2(5)	2.35(1)	2.6(5)	2.65(2)	7.8	1 (fix)	3.99 (fix)	1.8*ssO1	2<k<10	1.1<R<4.0
450	4.2(6)	2.35(2)	3.1(1)	2.64(2)	7.3	1 (fix)	3.99 (fix)	1.9*ssO1	2<k<10	1.1<R<4.0

NOTE: applied constraints, i.e. $n_{Cl} @ T_i \leq n_{Cl} @ T_{i+1}$, and $n_{tot} @ T_i \geq n_{tot} @ T_{i+1}$.

*fix based on everyone else's determination of $n=8$ at ambient conditions, use this to determine S_0^2 .

**Data at 450°C is noisier than at 400°C; equally good fits are obtained up to $n_{Cl} = 3.3$.

Table 6. Gibbs free energy of reactions and stepwise stability constants of Y(III)-Cl association reactions, calculated from box 1 and box 3. $\Delta_r G$, Gibbs free energy for the reaction; $\Delta_r G^{\ominus,c}$, Gibbs free energy at infinite dilution; $\Delta_r G^{\ominus}$, Gibbs free energy with concentration and activity correction.

Reaction in box 1	T(°C), P(bar)	$\Delta_r G$ (kJ/mol)	$\Delta_r G^{\ominus,c}$ (kJ/mol)	$\Delta_r G^{\ominus}$ (kJ/mol)	$\log K^{\ominus}$
$Y^{3+} + Cl^- = YCl^{2+}$	200, 800	-3.92 ± 8.85	3.60	-8.27	0.91 ± 0.98
	350, 800	-24.35 ± 10.42	-14.46	-55.96	4.69 ± 0.87
	500, 1000	-50.34 ± 11.60	-38.06	-146.44	9.89 ± 0.78
$YCl^{2+} + Cl^- = YCl_2^+$	200, 800	1.84 ± 9.21	7.30	-1.53	0.17 ± 1.01
	350, 800	-26.48 ± 10.11	-19.30	-45.33	3.80 ± 0.87
	500, 1000	-37.31 ± 11.37	-28.40	-93.27	6.30 ± 0.77
Reaction in box 3					
$Y^{3+} + Cl^- = YCl^{2+}$	200, 800	-12.05 ± 8.01	-2.21	-9.95	1.01 ± 0.89
	350, 800	-28.22 ± 9.48	-15.26	-58.19	4.88 ± 0.79
	500, 1000	-36.07 ± 10.06	-19.98	-142.58	9.63 ± 0.68
$YCl^{2+} + Cl^- = YCl_2^+$	200, 800	-2.12 ± 8.74	6.73	0.64	-0.07 ± 0.96
	350, 800	-26.23 ± 10.72	-14.58	-42.32	3.55 ± 0.90
	500, 1000	-30.62 ± 9.86	-16.16	-92.95	6.28 ± 0.67
$YCl_2^+ + Cl^- = YCl_3(aq)$	200, 800	2.70 ± 9.69	10.21	8.12	-0.90 ± 1.07
	350, 800	-12.19 ± 9.36	-2.30	-16.15	1.35 ± 0.78
	500, 1000	-33.39 ± 9.97	-21.12	-61.74	4.17 ± 0.67
$YCl_3(aq) + Cl^- = YCl_4^-$	200, 800	1.29 ± 8.46	6.75	6.75	-0.74 ± 0.93
	350, 800	-5.76 ± 9.15	1.42	1.42	-0.12 ± 0.77
	500, 1000	-17.84 ± 10.63	-8.93	-8.93	0.60 ± 0.72

Table 7. Cumulative formation constant of Y(III)-Cl complexes.

Reaction	T(°C), P(bar)	logβ⁰	error
Y³⁺ + Cl⁻ = YCl²⁺	200, 800	0.96	0.66
	350, 800	4.79	0.59
	500, 1000	9.76	0.50
Y³⁺ + 2 Cl⁻ = YCl₂⁺	200, 800	1.01	0.95
	350, 800	8.47	0.86
	500, 1000	16.05	0.71
Y³⁺ + 3 Cl⁻ = YCl₃(aq)	200, 800	0.11	1.43
	350, 800	9.82	1.16
	500, 1000	20.22	0.98
YCl₃(aq) + Cl⁻ = YCl₄⁻	200, 800	-0.63	1.71
	350, 800	9.70	1.39
	500, 1000	20.82	1.22

Table 8. Modified Ryzhenko–Bryzgalin (MRB) (Ryzhenko et al., 1985; Shvarov and Bastrakov, 1999) equation of state parameters for the Y(III)-Cl complexes obtained via fitting of $\log K^\ominus$ values from MD simulations. *The B parameters were set to zero, except for the association reaction of $Y^{3+} + Cl^- = YCl^{2+}$, whose $\log \beta_{(25^\circ C, 1 \text{ bar})}^\ominus$ was chosen from Luo & Byrne (2001). [you have K is italics sometimes...]

reaction	$\log \beta_{(25^\circ C, 1 \text{ bar})}^\ominus$	A_{zz}	B_{zz}^*
$Y^{3+} + Cl^- = YCl^{2+}$	0.65	3.357	- 1428
$Y^{3+} + 2 Cl^- = YCl_2^+$	-8.314	3.047	-
$Y^{3+} + 3 Cl^- = YCl_3(aq)$	-12.914	3.97	-
$Y^{3+} + 4 Cl^- = YCl_4^+$	-14.800	4.176	-

Table 9. the logarithm of formation constant Y(III)-Cl complexes calculated using MRB parameters in Table 8. *, values are used in discussion part.

T (° C)	P _{Sat}	800 bar	1000 bar	2000 bar	5000 bar	P _{Sat}	800 bar	1000 bar	2000 bar	5000 bar
$Y^{3+} + Cl^- = YCl^{2+}$						$Y^{3+} + 2 Cl^- = YCl_2^+$				
25	0.65	1.04	1.13	1.51	2.26	-8.31	-9.14	-9.33	-10.14	-11.74
50	0.22	0.48	0.54	0.81	1.36	-6.58	-7.34	-7.51	-8.27	-9.86
100	0.02	0.13	0.16	0.27	0.51	-3.38	-4.14	-4.30	-5.03	-6.61
150	0.43	0.43	0.43	0.44	0.45	-0.50	-1.37	-1.55	-2.33	-3.96
200	1.24	1.12	1.09	0.99	0.80	2.21*	1.12	0.90	0.01	-1.76
250	2.37*	2.06	2.00	1.78	1.37	4.96*	3.46	3.18	2.09	0.12
300	3.87*	3.21	3.10	2.71	2.07	8.11	5.78	5.40	4.02	1.76
350	6.21*	4.58	4.39	3.75	2.84	12.90	8.24	7.68	5.87	3.24
400		6.26	5.90	4.89	3.64		11.05	10.15	7.68	4.60
450		8.39	7.69	6.13	4.48		14.48	12.94	9.49	5.86
500		11.14	9.85	7.45	5.33		18.77	16.16	11.32	7.05
$Y^{3+} + 3 Cl^- = YCl_3(aq)$						$Y^{3+} + 4 Cl^- = YCl_4^+$				
25	-12.91	-14.00	-14.24	-15.30	-17.38	-14.80	-15.94	-16.19	-17.31	-19.50
50	-10.49	-11.48	-11.71	-12.69	-14.77	-12.16	-13.20	-13.43	-14.47	-16.66
100	-6.07	-7.06	-7.27	-8.22	-10.28	-7.35	-8.39	-8.62	-9.62	-11.78
150	-2.12	-3.25	-3.49	-4.50	-6.62	-3.09	-4.27	-4.52	-5.59	-7.82
200	1.56	0.14	-0.14	-1.30	-3.60	0.88	-0.61	-0.91	-2.14	-4.55
250	5.27*	3.32	2.95	1.54	-1.03	4.85	2.79	2.41	0.93	-1.78
300	9.49	6.45	5.95	4.16	1.22	9.35	6.15	5.62	3.74	0.65
350	15.82	9.74	9.01	6.65	3.23	16.06	9.67	8.89	6.41	2.81
400		13.47	12.30	9.08	5.07		13.63	12.40	9.01	4.79
450		18.01	16.00	11.50	6.77		18.45	16.33	11.60	6.62
500		23.65	20.25	13.94	8.38		24.41	20.84	14.20	8.35

Figure 1. Snapshots of Y(III)-Cl complexes at different T, P from MD simulations. (a-d) are configurations obtained from simulations of box 1 under condition of 25 °C 1 bar; 200 °C, 800 bar; 350 °C, 800 bar; 500 °C, 1000 bar, respectively; (f) is the configuration derived at 500 °C and 1000 bar in box 3.

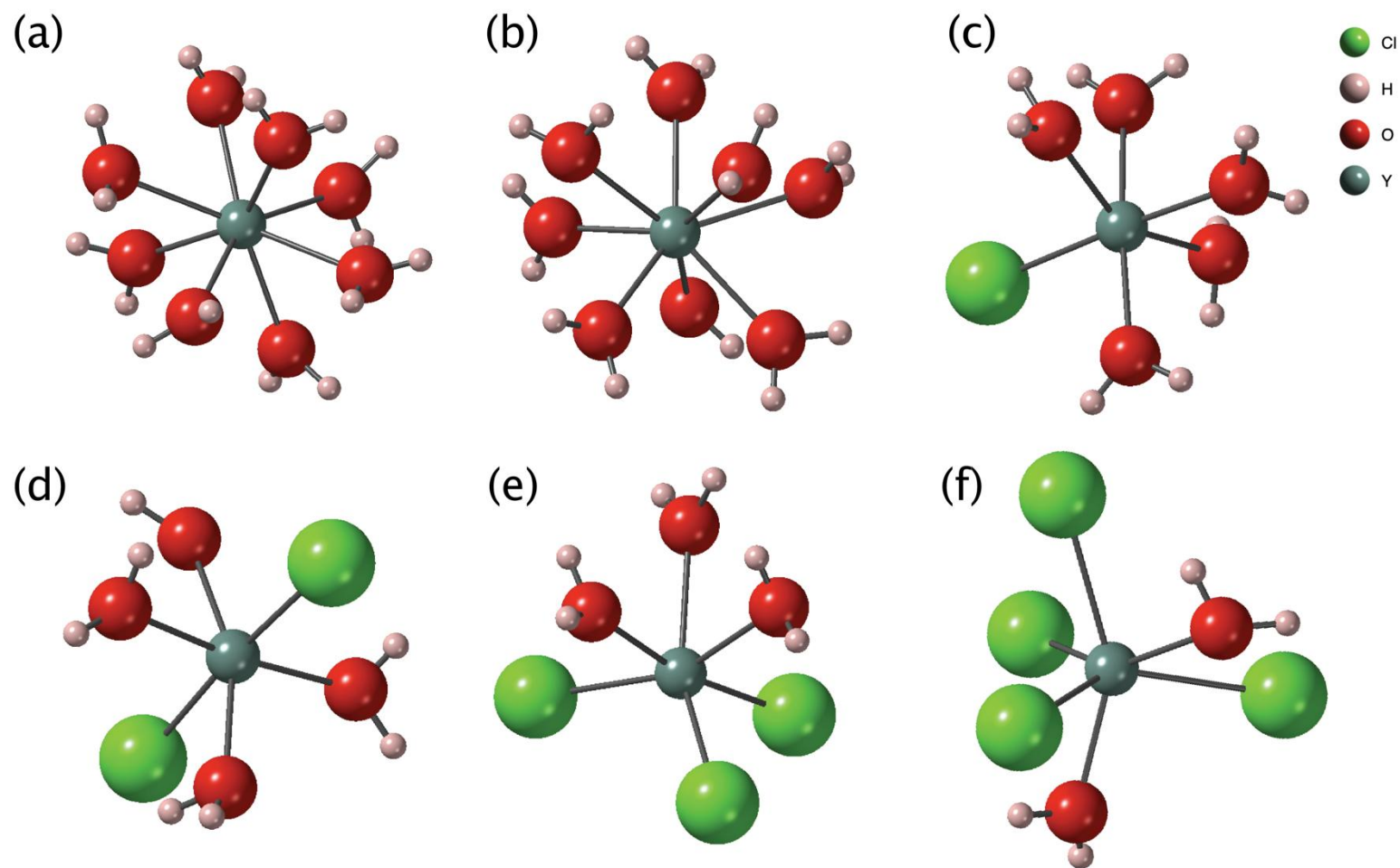


Figure 2, Radial distribution functions (RDF, left axes, solid lines) and their integrals (coordination number, CN, right axes, dash lines) of Y(III)-Cl and Y(III)-O (x-axis) in the four boxes of simulations, (a), (b), box 1; (c),(d), box 2; (e), (f), box 3; (g), (h), box 4.

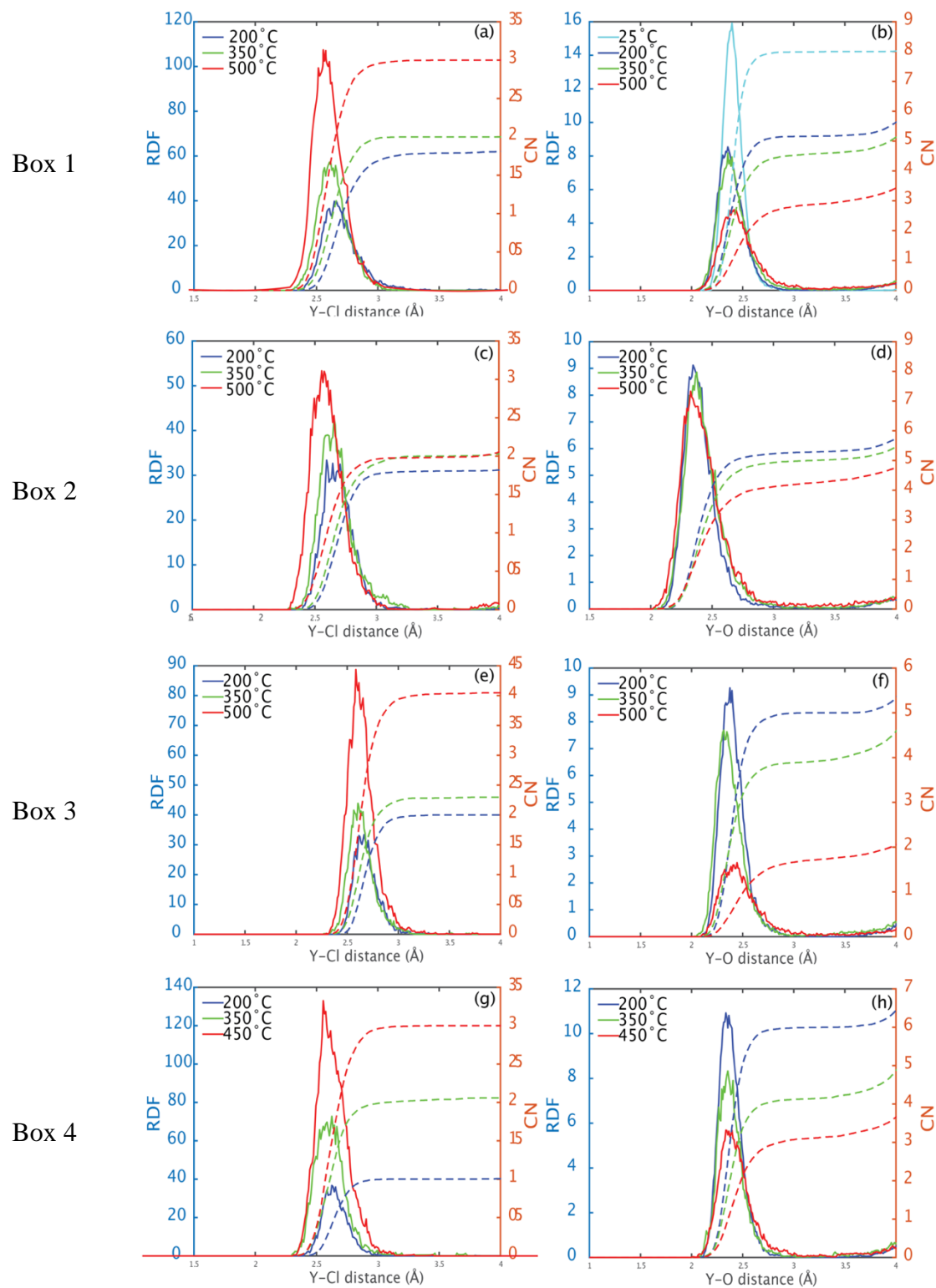


Figure 3. Y(III)-Cl distances over simulation time of (a), job 1a(2) and (b), 1b.

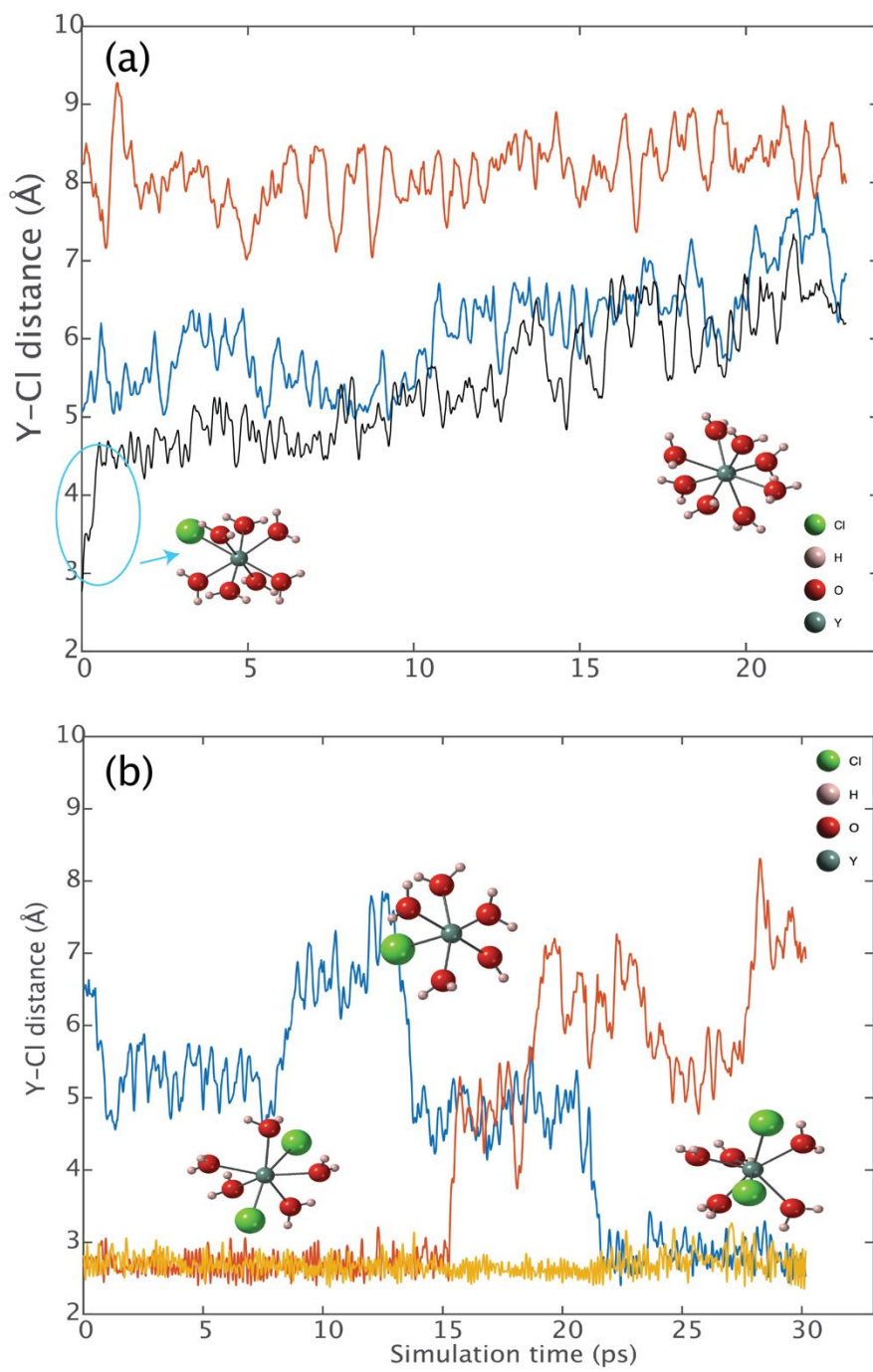


Figure 4. XANES spectra on the solution with highest chlorine concentration (Sol5; 10 m LiCl) as a function of temperature.

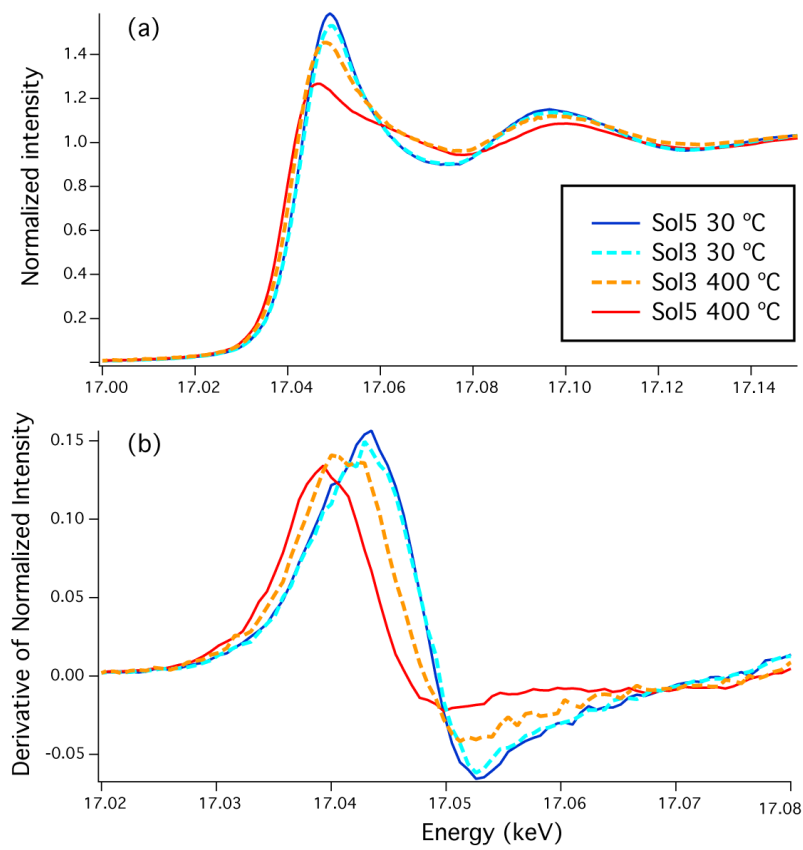


Figure 5. EXAFS data (solid lines) and shell-by-shell fits (dotted lines).

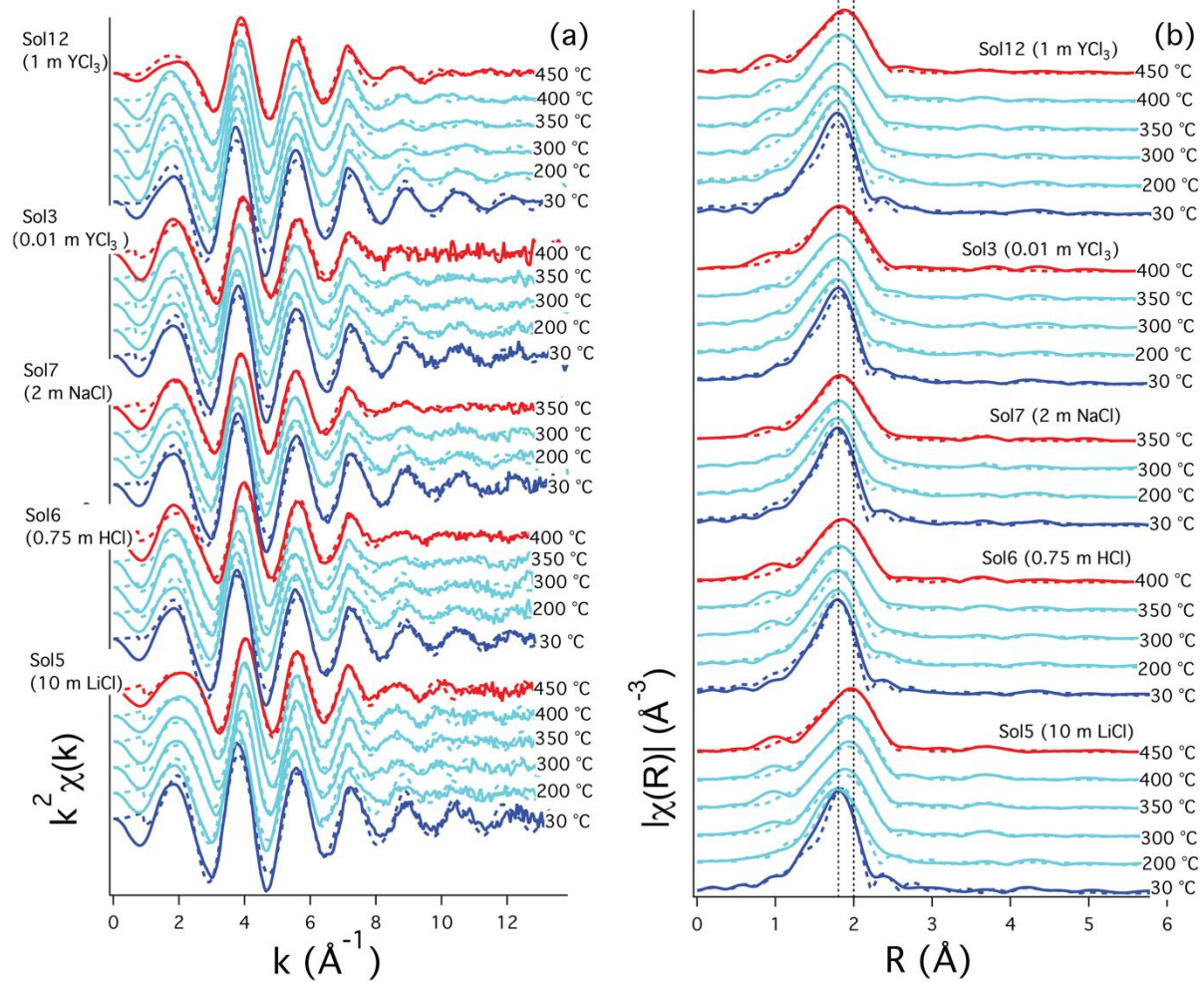


Figure 6. Experimental and calculated XANES spectra: The Yttrium K-edge (17,038 eV) was subtracted from the experimental energy to be compared with the FDMNES spectra. (a) crystalline $Y_2O_3(s)$; the crystal structure used for FDMNES is from Santos et al. (2005); (b) antiprism $[YO_8]$ moiety with different bond distance (r) and angle between Y-O bond and z axis (angle θ); (c) triangular prismatic $[YCl_nO_{6-n}]$ ($n \leq 3$) moiety with bond distances $r_{Y-O} = 2.37 \text{ \AA}$ and $r_{Y-Cl} = 2.65 \text{ \AA}$, compared to selected Y-O-Cl complexes from the MD simulations; and (d) the average of calculated spectra from MD boxes compared to experimental spectra over a range of conditions.

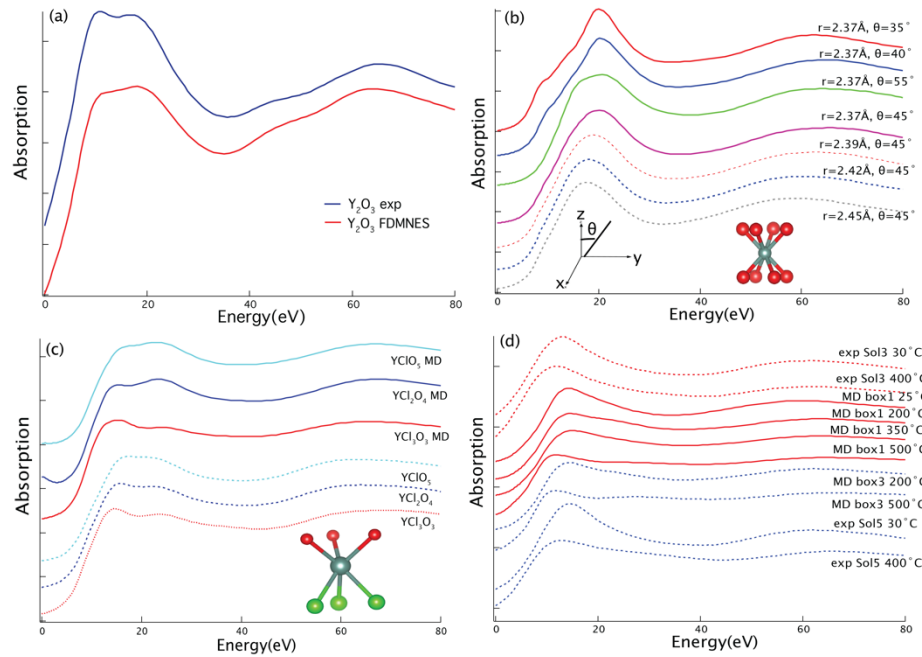


Figure 7. Constraint mean force and free energy surface of dissociation reaction of YCl^{2+} at 200 °C, 800 bar.

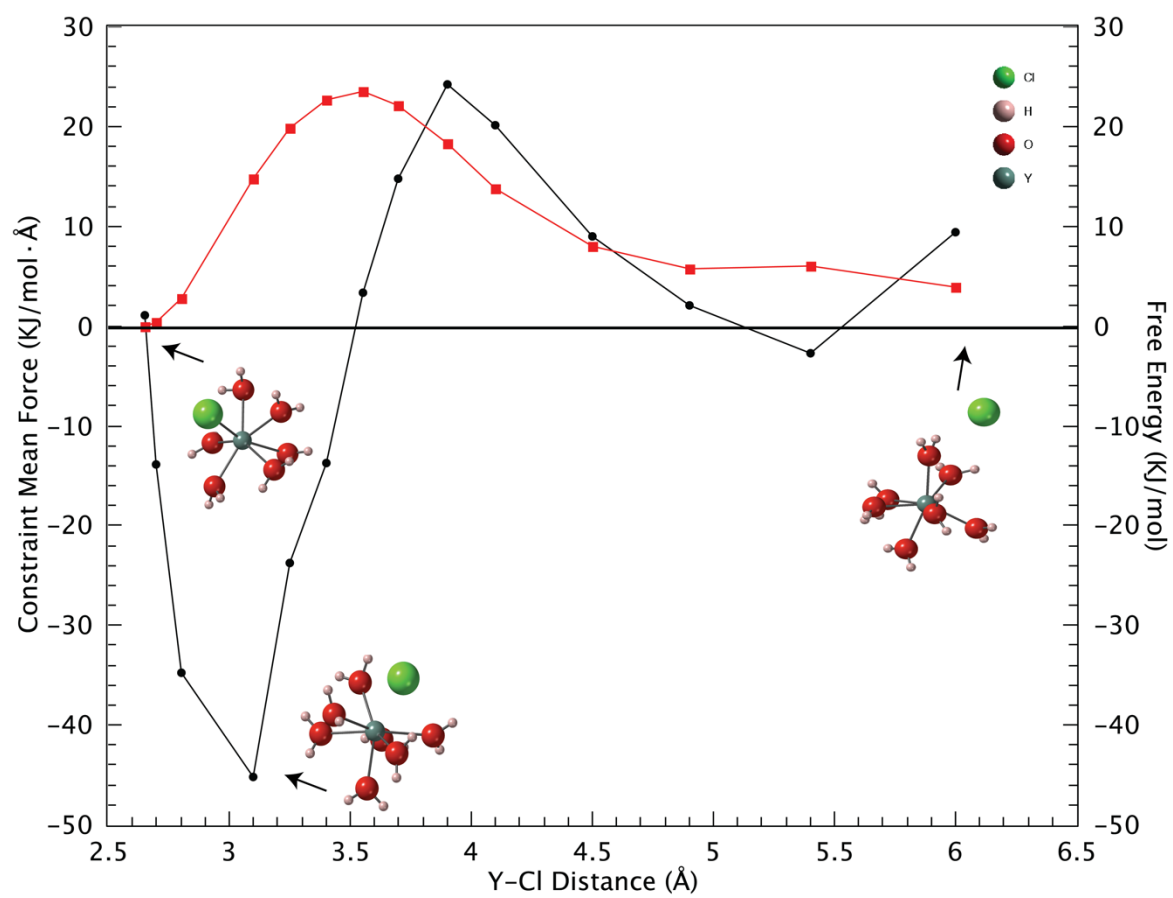


Figure 8. Free energy surface of Y(III)-Cl complexes at 200 °C, 800 bar; 350 °C, 800 bar; 500 °C, 1000 bar from thermodynamic integration calculation in box 1 (a - c) and box 3(d - f).

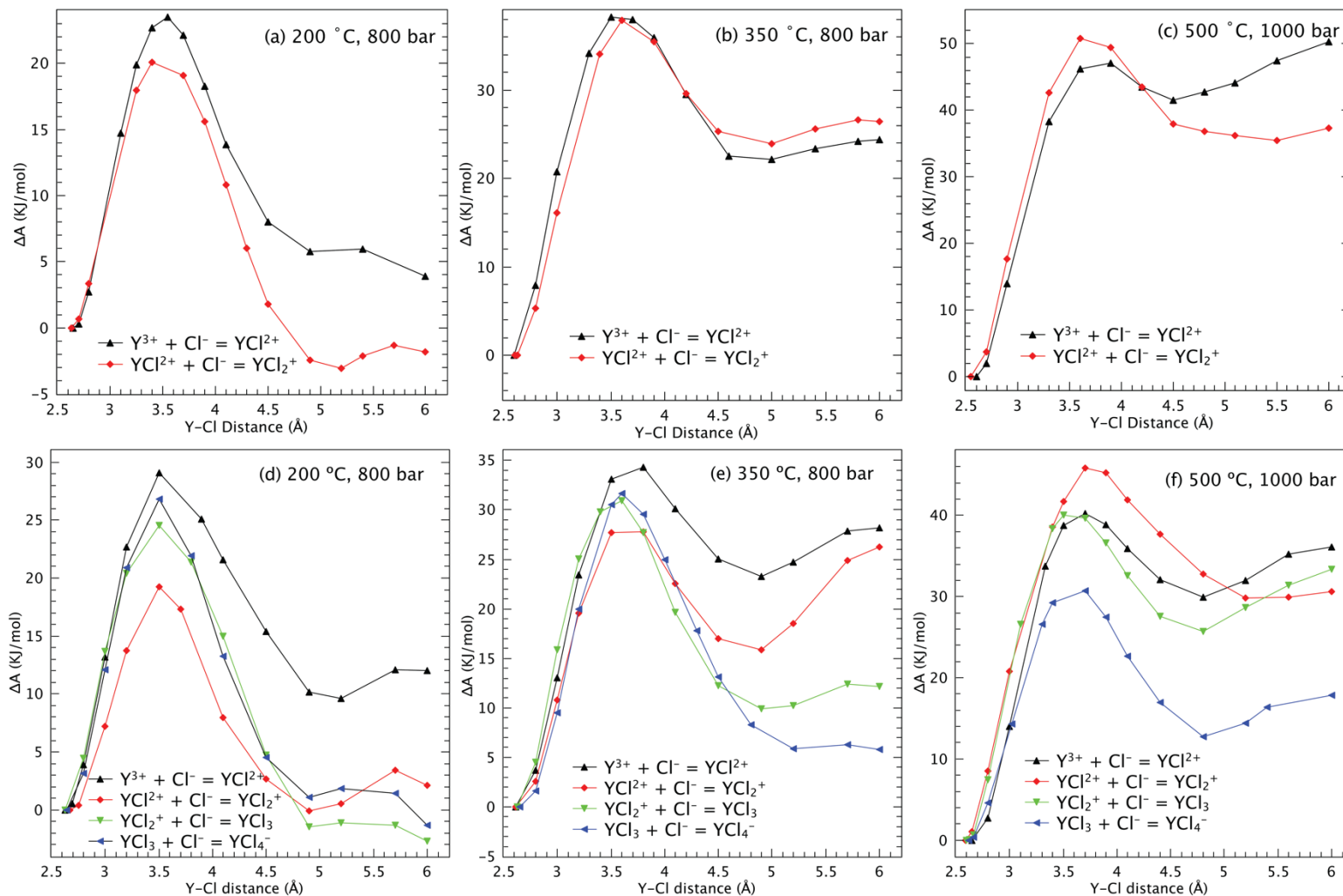


Figure 9. The stepwise $\log K^\ominus$ of association of Y(III)-Cl complexes at 200 °C, 800 bar; 350 °C, 800 bar; 500 °C, 1000 bar and within error bar, and the average value of the two boxes.

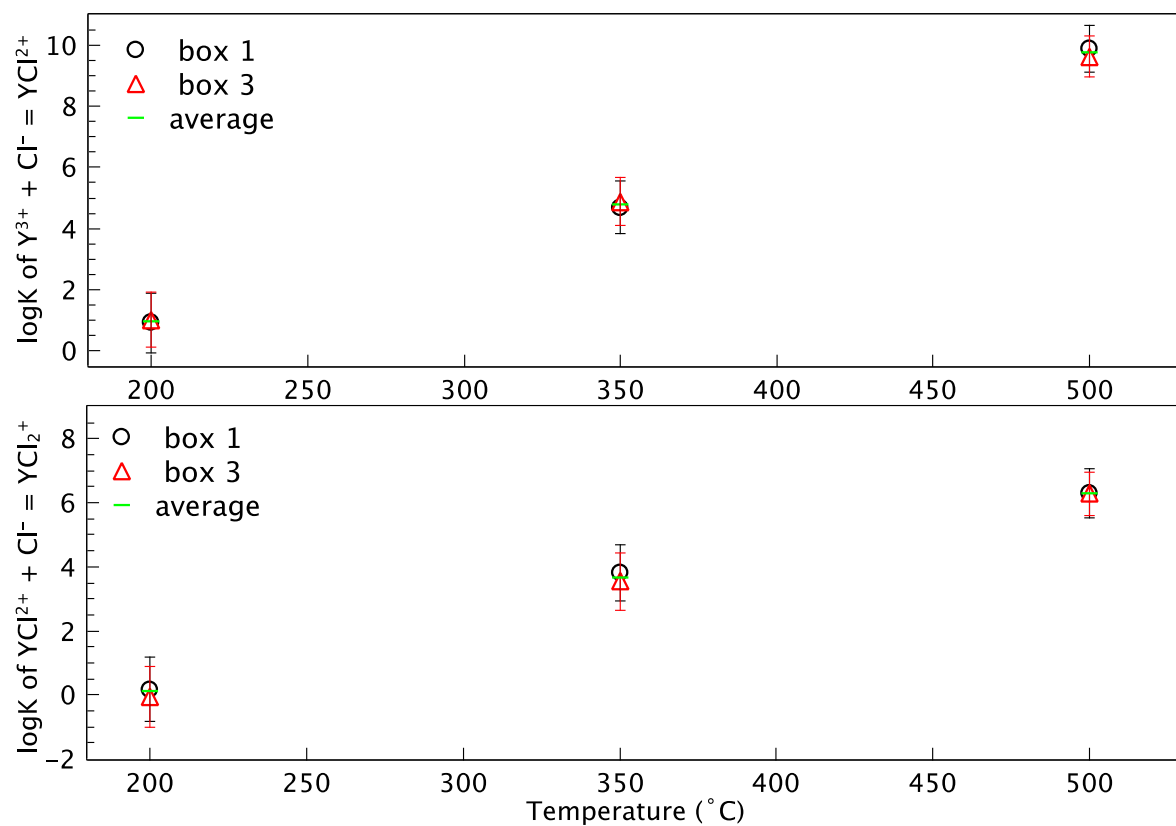


Figure 11. The distribution of Y(III) chloride species at 200 °C, 800 bar; 350 °C, 800 bar; 500 °C, 1000 bar, and pH =3 and 0.3. The solution starts with 0.01 m of Y^{3+} and 0.03 m Cl^- . Dashed lines are average coordination numbers (right axis) from the modelled distribution of species, orange dots are average coordination numbers from XAS Sol 3 at $n(Cl^-) = 0.003$ m (a); Sol 7 at $n(Cl^-) = 2.5$ m and (b,c) Sol 5 at $n(Cl^-) = 5$ m (b,d,f). Errors bars corresponding to ± 0.2 Cl^- ions were added for XAS solutions where this number was fixed, based on fits using different fixed numbers.

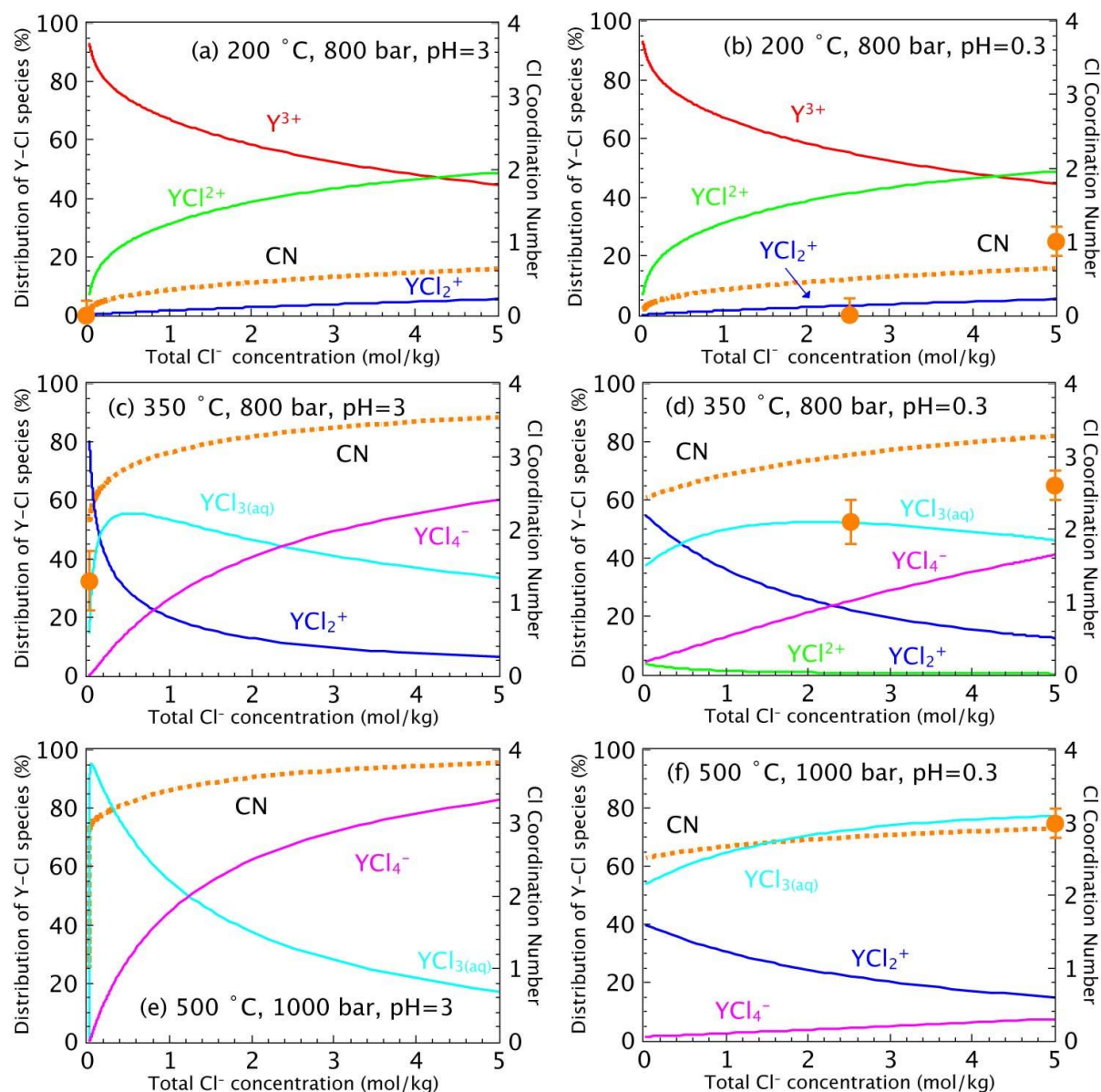


Figure 12. Comparison of the formation constants of REECl_2^{2+} and REECl_2^+ , measured from solubility experiment from Migdisov (2009); ∇ , predicted from HKF parameters from Migdisov (2016); \times , predicted from MRB model in this study.

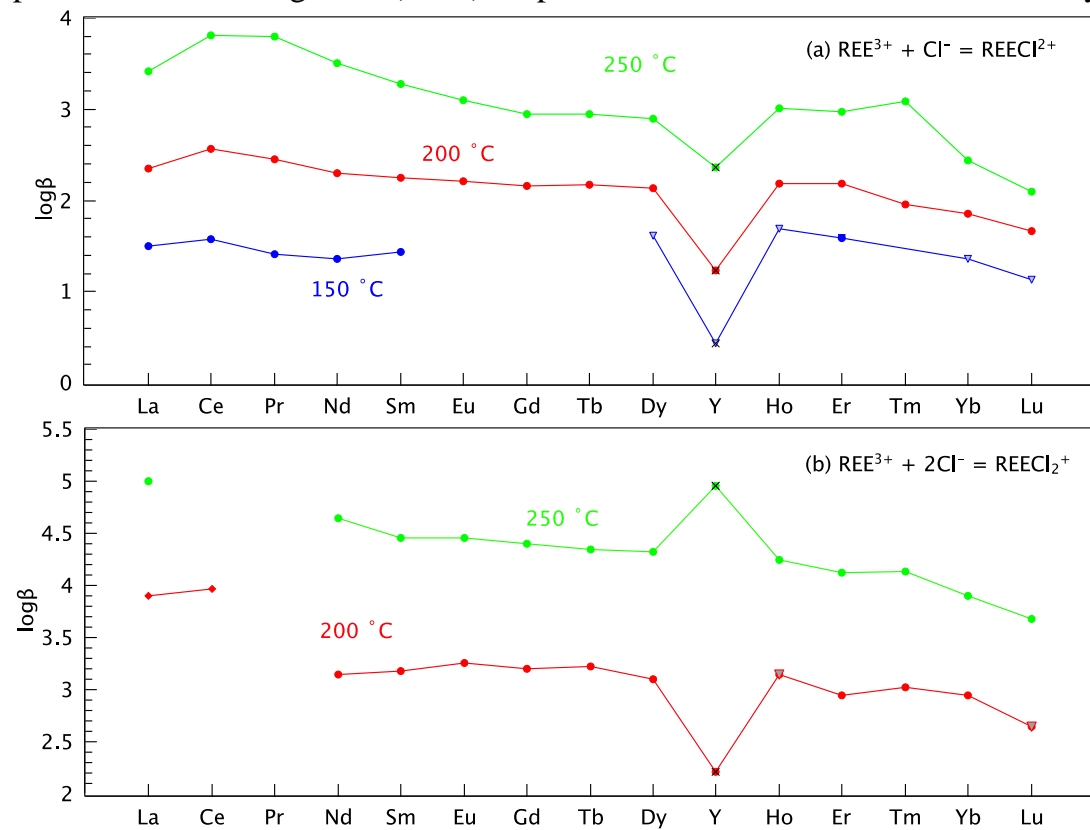


Figure 13. Ho-Cl/Y-Cl species with Cl concentration at 250 °C P_{sat} . The solution starts with 0.001 m Y/Ho and 0.003 m Cl^- , pH=4.

