

1 PyXRD v0.6.7: a free and open source program to quantify disordered
2 phyllosilicates using multispecimen X-ray diffraction profile fitting.

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12 **Abstract**

13 This paper presents a free and open-source program called PyXRD (short for Python X-
14 ray diffraction) to improve the quantification of complex, poly-phasic mixed-layer
15 phyllosilicate assemblages. The validity of the program was checked by comparing its
16 output with Sybilla v2.2.2, which shares the same mathematical formalism. The novelty of
17 this program is the ab initio incorporation of the multispecimen method, making it possible
18 to share phases and (a selection of) their parameters across multiple specimens. PyXRD
19 thus allows modelling multiple specimens side by side, and this approach speeds up the
20 manual refinement process significantly. To check the hypothesis that this multispecimen
21 set-up – as it effectively reduces the number of parameters and increases the number of
22 observations – can also improve automatic parameter refinements, we calculated X-ray
23 diffraction patterns for four theoretical mineral assemblages. These patterns were then
24 used as input for one refinement employing the multispecimen set-up and one employing
25 the single-pattern set-ups. For all of the assemblages, PyXRD was able to reproduce or
26 approximate the input parameters with the multispecimen approach. Diverging solutions
27 only occurred in single-pattern set-ups which do not contain enough information to discern
28 all minerals present (e.g. patterns of heated samples). Assuming a correct qualitative
29 interpretation was made and a single pattern exists in which all phases are sufficiently
30 discernible, the obtained results indicate a good quantification can often be obtained with
31 just that pattern. However, these results from theoretical experiments cannot automatically
32 be extrapolated to all real-life experiments. In any case, PyXRD has proven to be useful
33 when X-ray diffraction patterns are modelled for complex mineral assemblages containing
34 mixed-layer phyllosilicates with a multispecimen approach.

35 **Keywords:** quantitative, XRD, clay mineralogy, PyXRD, multispecimen

36 **1. Introduction**

37 Clay minerals (i.e. phyllosilicates) are among the most difficult minerals to study in detail
38 due to their inherent chemical and structural variability (Środoń, 2006; Velde and Meunier,
39 2008; Hubert et al., 2012). Nonetheless, these minerals are one of the most abundant
40 constituents of the Earth's upper crust, and have an important influence on various
41 physical (e.g. plasticity, shear strength, porosity) and chemical (e.g. buffering and
42 exchange capacities, pH, electrical conductivity) properties (Agbenin and Tiessen, 1995;
43 Vernik and Liu, 1997; Righi et al., 1999; Wen and Aydin, 2003; Lado and Ben-Hur, 2004;
44 Caner et al., 2010). Phyllosilicates are also very reactive phases responding quickly to
45 changes in their environment (Pai et al., 2004; Meunier, 2007; Velde and Meunier, 2008;
46 Cornelis et al., 2014).

47 Therefore, quantitative information on the mineralogical composition of clay-bearing
48 samples is an important step in characterizing and understanding them. Different
49 techniques can be used to quantify clay minerals, but those using X-ray diffraction are the
50 most abundant and have proven to be the most reliable (Plançon, 1981; Reynolds Jr.,
51 1985; Drits and Tchoubar, 1990; Righi et al., 1999; Sakharov et al., 1999a; Środoń, 2006;
52 Hubert et al., 2009, 2012; Ufer et al., 2012a; b). Models calculating X-ray diffraction
53 patterns usually provide the highest level of detail because the input for such models can
54 be considered an approximation of the real structure of the minerals (e.g. layer structures,
55 composition, stacking parameters, interlayer composition, orientation). As such, this
56 approach does not only yield quantitative data, but also structural and compositional
57 information. However, this also means a large number of variables are involved, some of
58 which are very difficult to predict or estimate in advance. In combination with the complex,
59 poly-phasic nature of many natural samples, it is a challenge to create software that allows
60 for the completely automated quantification of clay minerals.

61 Two complementary methods exist to analyse clay minerals using X-ray diffraction. One
62 uses powder samples, for which the orientation of crystallites is considered to be
63 approximately random, and the other uses oriented samples, in which the orientation of
64 crystallites occurs mainly along a plane of preferred orientation. Originally, powder X-ray
65 diffraction was and still is used to determine crystal structures for unknown phases (not
66 just phyllosilicates), which then developed into quantitative analysis. However, for
67 disordered structures like mixed-layered clay minerals, powder patterns are often difficult
68 to interpret. In such cases oriented patterns can be used to focus on the stacking
69 (dis)order along the c^* axis. Since the 1970's, several computer programs have been
70 developed to calculate X-ray diffraction patterns for (disordered) clay minerals (Kakinoki &
71 Komura 1965; Reynolds, 1967; Ergun, 1970; Drits & Sakharov, 1976; Plançon, 1981).
72 Examples of commonly used programs are the NEWMOD[®]-family (Reynolds Jr., 1985;
73 Pevear and Schuette, 1993; Reynolds Jr. and Reynolds III, 1996; Yuan and Bish, 2010),
74 MLM2C/3C and derivatives (Plançon and Drits, 2000), Sybilla (Aplin et al., 2006; McCarty,
75 2015), DIFFaX (Treacy et al., 1991) and BGMN (Ufer et al., 2012a; b). Some of these
76 programs (e.g. DIFFaX, BGMN, Wildfire, Sybilla 3D) are able to calculate X-ray diffraction
77 patterns for random powder diffraction patterns, while others (NEWMOD[®], MLM2C,
78 MLM3C, Sybilla) focus only on calculating one-dimensional (00l) patterns.

79 Another aspect to consider is the ability of these programs to automatically refine
80 parameters. For instance, the last version of NEWMOD[®] uses a simple linear least-
81 squares algorithm, Sybilla makes use of a genetic algorithm, and BGMN has a custom
82 least-squares algorithm. In essence, all of these algorithms try to find a solution by
83 minimizing a target function, usually a measure for the difference between the calculated
84 and observed data. This difference is usually defined as the sum of the squares of the
85 errors or as the pattern's R_p factor (Toby, 2006). A linear or ordinary least-squares

86 algorithm works well when there is a well-defined global minimum and the target function
87 is relatively smooth. However, for more complex cases this is often not the case, and as a
88 result an ordinary least-squares might not converge at all. Algorithms using a more
89 stochastic approach, like genetic algorithms, can partly overcome problems related to
90 target function smoothness or poorly defined minima (also see section 2.4). Nonetheless,
91 any algorithm will require some guidance e.g. by not releasing all parameters for automatic
92 refinement at once, by adjusting some parameters manually, by setting upper and lower
93 limits or by choosing starting values close enough to the actual solution. The reason is that
94 models describing X-ray diffraction by disordered layered minerals can not always be
95 constrained adequately, and a successful quantification is still very dependent on the skill
96 of the individual modeller. As a result, most published quantifications of complex mixed-
97 layer assemblages employ a time-consuming trial-and-error approach at some point in the
98 modelling process.

99 Several authors used a 'multispecimen approach' to further constrain their models (Drits,
100 1997; Sakharov et al., 1999a; b; Hubert et al., 2012 and references therein). This
101 approach involves recording multiple specimens (e.g. air-dried, glycolated, heat
102 treatments) of the same sample and creating a structural model that can explain the
103 observed features for all specimens. The reason for doing this is that swelling layers (like
104 smectite and vermiculite layers) will expand or contract in response to these treatments.
105 The level of expansion or contraction can be related to layer charges, and helps in
106 discerning the different swelling phases present and understanding their stacking
107 (dis)order (Ferrage et al. 2005a,b; Michot et al., 2005; Ferrage et al. 2007; Dazas et al.,
108 2015). In short, this approach allows to determine the structure and type of (mixed-layer)
109 clay minerals present in the sample with higher certainty. However, today not a single
110 program allows for a side-by-side calculation of these patterns. Because of this, modellers

111 are still forced to refine their model parameters on one specimen and then check if the
112 solution also explains the other observations. As long as a manual trial-and-error
113 refinement process is used, this does not pose too many practical problems aside from the
114 time needed. In theory however, a program able to integrate all the observations and
115 calculate patterns for them could lead to better automatic parameter refinements, a
116 hypothesis tested in this paper using theoretical assemblages.

117 The program presented in this paper, called PyXRD (short for Python X-ray Diffraction),
118 was designed with this multispecimen approach in mind. It (selectively) shares phase
119 parameters across specimens and keeps phase quantities identical in each specimen,
120 thus reducing the number of parameters while at the same time increasing the number of
121 observations. Other design goals for PyXRD were (i) to have an easy-to-use interface, (ii)
122 to be an open program allowing as many aspects of the input to be changed as possible,
123 (iii) to provide a means for automatic parameter refinement, and (iv) to provide an open-
124 source program for others, allowing them to use the software freely and make
125 improvements where they see fit.

126 This paper illustrates the general structure of this program and presents the results from a
127 comparison between PyXRD and Sybilla v2.2.2 and between automatic parameter
128 refinements for several theoretical mineral assemblages, with and without the use of the
129 multispecimen approach. The software manual contains more detailed information about
130 the numerical solutions used for calculating the X-ray diffraction patterns and a guided
131 example on how to create projects using the graphical user interface (GUI).

132 **2. Materials and methods**

133 **2.1 Model implementation and licence**

134 PyXRD is written in Python 2.7 and uses a number of open-source third-party modules.

135 The GUI utilizes PyGTK as widget toolkit and has an internal model-view-controller
136 framework. To improve calculation speed, PyXRD makes use of the NumPy and SciPy
137 libraries. NumPy provides multidimensional array objects and many related routines for
138 manipulating them, while SciPy provides more complex mathematical and scientific
139 algorithms built on top of NumPy (Jones et al., 2001; van der Walt et al., 2011). The
140 Matplotlib library is used for plotting patterns and data (Hunter, 2007). Finally, the
141 Distributed Evolutionary Algorithms for Python (DEAP) library is used to harness to power
142 of evolutionary algorithms to automatically refine parameters (Fortin et al., 2012).
143 PyXRD is released under a BSD licence, except for the mvc module which, as it is a
144 derived work from the gtkmvc project, is licensed as GNU LGPL v2.

145 **2.2 Program data structure**

146 PyXRD is implemented according to a model-view-controller (mvc) paradigm separating
147 data and calculations from GUI-related aspects. In the following section, an overview is
148 given of the most important objects found in the data layer and their associations. More
149 details can also be found in the manual and the source code documentation.

150 **Project object**

151 The user interface of PyXRD can create (or load) a single *Project* object. It is a container
152 object grouping lists of *AtomType*, *Phase*, *Specimen* and *Mixture* objects together. These
153 are the four top-level objects which are used to calculate X-ray diffraction patterns. Their
154 associations are shown schematically in figure 1. The purpose of each of them will be
155 explained in more detail below.

156 **AtomType object**

157 The *AtomType* object is the most basic building block. This object bundles all the physical

158 constants (e.g. charge, atomic weight, scattering factors) for a single ion (e.g. Fe²⁺, Fe³⁺)
159 or for a molecule (e.g. H₂O and glycol) small enough to be considered having a spherical
160 electron cloud. When a new project is created, a default list of these *AtomType* objects is
161 loaded, using the atomic scattering factors as published by Waasmaier and Kirfel (1995).

162 **Phase and Component objects**

163 *Phase* objects contain all the information needed to calculate a one-dimensional X-ray
164 diffraction pattern of a (mixed-layer) mineral. A *Phase* combines (i) a *Probability* object, (ii)
165 an object describing the coherent scattering domain size (*CSDS*), and (iii) one or more
166 *Component* objects which contain information about the structure of the different types of
167 layers in the *Phase*.

168 The *Probability* object describes how these layers are stacked by means of Markovian
169 statistics and the Reichweite concept (Drits and Tchoubar, 1990). Currently PyXRD has
170 implemented probability models for R values ranging from 0 to 3. For each combination of
171 Reichweite and number of components there are a number of independent parameters
172 required to calculate the remaining parameters, which describe the stacking order or
173 disorder. The values of these independent parameters can be based on another phase
174 with the same combination of Reichweite and number of components. For example, this
175 means it is possible to share the illite content in an illite/smectite mixed layer across its AD
176 and EG phase, but have different weight fractions (or junction probabilities) for the different
177 types of smectite in those phases. For a complete explanation on how these calculations
178 work and which parameters were chosen to be independent we refer to the manual.

179 The *CSDS* object describes what type of coherent scattering domain size distribution
180 should be used and contains the necessary parameter values to describe this distribution
181 (e.g. average *CSDS*). Two types of *CSDS* distributions are currently implemented: a
182 generic log-normal distribution and a log-normal distribution in which the distribution

183 constants published in Drits et al. (1997) are employed and the average CSDS is the only
184 remaining unknown variable. Each phase also has a σ^* factor which makes it possible to
185 correct for incomplete preferred orientation (Reynolds Jr., 1986; Dohrmann et al., 2009).
186 The *Component* object describes the size, structure, composition and (variation in) basal
187 spacing of a single layer type in that phase. A *Component* contains two lists that combine
188 an *AtomType* from the project with its (projected) coordinate along the c*-axis (also known
189 as the z coordinate) and the number of projected ions of that type at that coordinate. The
190 first list involves atoms in the silicate lattice, while the other list describes the (variable)
191 interlayer space. With this approach, the silicate structure can be shared between different
192 phases (e.g. AD and EG states), while the interlayer contents may still be different.

193 **Specimen objects**

194 *Specimen* objects provide all the information regarding the experimental data (the actual
195 measurements, sample size, etc.) and the *Goniometer* set-up (radius, slit sizes, etc.). They
196 do not hold a direct reference to phases, but are linked with them through *Mixture* objects.

197 **Mixture objects**

198 *Mixture* objects are the starting point for the actual calculations as they link phases and
199 specimens together. In the user interface, a table can be created by adding just as many
200 rows as there are *Phases* and just as many columns as there are *Specimens*. In the
201 column headers, there are slots where the user can select the *Specimen*. Similarly, the
202 user can select the corresponding *Phase* in each cell of the table. This enables the user to
203 select different states of smectite for an AD and an EG *Specimen* (see figure 2 for a
204 screenshot of the GUI), while keeping unaffected *Phases*, (e.g. kaolinites, micas and
205 chlorites) unchanged.

206 Once a *Mixture* is created in this way, a number of parameters are available for automatic

207 refinement (e.g. weight fractions from the *Probability* object, the average CSDS, etc.). In a
208 refinement dialog, the user can select which parameters s/he would like to improve and
209 the minimum and maximum values between which the ideal value should lie. A number of
210 different refinement methods are also available - some of them more complex or
211 specialized than others. Yet, as a complete description of all methods is beyond the scope
212 of this article, only the algorithm used for the refinements will be explained in detail below.

213 **2.3 Numerical calculations**

214 The X-ray diffraction patterns are calculated using the matrix formalism, for which a very
215 good summary can be found in Drits and Tchoubar (1990). Later developments
216 incorporated can be found in Drits et al. (1997) and Plançon (2001). Since the complete
217 mathematical deduction followed for PyXRD is rather long, in itself does not contain new
218 developments, and is not the aim of this paper it is not included here. However, an
219 overview of the mathematical deductions and calculations, as they are implemented in the
220 'calculations' module, can be found in the online manual
221 (<http://users.ugent.be/~madumon/pyxrd/Manual.pdf>) or in the manual included in this
222 article's supplement.

223 To improve calculation speed, programs can make use of multi-threading, spreading the
224 load from the different threads evenly over the different cores in a multi-core CPU.
225 However, multi-threading is not very effective in Python because of the Global Interpreter
226 Lock (GIL). This lock can only be obtained by a single active thread, while the others have
227 to wait for it to be released again. So instead of multi-threading, PyXRD uses multi-
228 processing, which creates a new python interpreter for each process, circumventing the
229 GIL problem. The downside is that processes, unlike threads, do not share memory.
230 Therefore, each process needs to be given all the data required to run the calculation. This
231 is achieved by isolating the calculation functions from objects and by extracting the

232 required data from the objects described in the previous section. As a result, the data
233 exchanged between processes is reduced to a minimum. This approach also makes it
234 possible to run PyXRD refinements effectively on high-performance computing (HPC)
235 clusters. The experiments presented in this paper were run on the HPC clusters of the
236 Stevin Supercomputer Infrastructure at Ghent University.

237 **2.4 Refinement algorithm**

238 PyXRD supports several refinement algorithms, but for more complex problems involving
239 several parameters, the genetic algorithms or evolutionary strategies are found to be most
240 reliable. PyXRD implements several evolutionary strategies, among which are a
241 Covariance Matrix Adaptation Evolutionary Strategy (Hansen and Ostermeier, 2001) and a
242 (multiple) Particle Swarm Optimization (Blackwell et al., 2008). While the Particle Swarm
243 Optimization is effective at searching the parameter space for minima, being able to
244 escape local minima easily, it can take a lot of function calls for it to converge. On the other
245 hand, the Covariance Matrix Adaptation Evolutionary Strategy is much more effective for
246 local searches, but does get stuck in local minima more easily. Therefore, PyXRD also
247 implements a Particle Swarm Covariance Matrix Adaptation Evolutionary Strategy
248 algorithm which extends the Covariance Matrix Adaptation Evolutionary Strategy with
249 collaborative concepts from a Particle Swarm Optimization (Muller et al., 2009), making it
250 the more robust choice. This Particle Swarm Covariance Matrix Adaptation Evolutionary
251 Strategy was also used for the experiments presented below.

252 **3. Results**

253 In the following sections PyXRD's output is compared with Sybilla's output. In the first
254 section, single phases are tested to check the implementation of the model. In the second
255 section a number of assemblages are tested to check if the obtained weight fractions are

256 correct. In the last section a comparison is made between single- versus multispecimen
257 refinements.

258 **3.1 Comparison between Sybilla and PyXRD results: calculated 00l** 259 **reflections for single discrete and mixed-layer phyllosilicates**

260 In total, 13 phases were tested. An overview of these phases with their most important
261 structural parameters are given in table 1. The original Sybilla projects, the produced
262 patterns and the PyXRD projects used can be found in this paper's supplement. All
263 patterns were calculated using a fixed σ^* value of 12, a sample length of 1.25 cm, a
264 goniometer radius of 17.3 cm, a divergence slit of 0.5° , Soller slits of 2.3° and an angular
265 range of 2° - 52° 2θ with 1000 steps (step size of 0.05° 2θ). The z^* coordinates of the atoms
266 were set to match with those in Sybilla, as were the scattering factors, the unit cell
267 dimension in the z direction, the octahedral iron content (for illite, chlorite and smectite
268 components), the interlayer water, glycol and cation contents (for smectite and illite
269 components) and the average coherent scattering domain size. The probability parameters
270 were entered as such that identical P and W matrices were obtained. For most of the
271 phases this meant identical parameters could be entered. Only for the R2 illite/smectite
272 with two components 2 additional parameters were entered in comparison with Sybilla,
273 which has a more restricted probability model for this combination of Reichweite and
274 components. These parameters are the junction probabilities P_{21} (fixed at 1.0 in Sybilla)
275 and P_{221} (fixed at 0.0 in Sybilla). A complete deduction on how the entered probabilities
276 and weight fractions are used to calculate the unknown weight and probability fractions is
277 present in the PyXRD manual. Sybilla uses scattering factors for the atoms in the silicate
278 lattice assuming 50% ionization, with the exception of Mg which is fully ionized (D.
279 McCarty, 2015). The scattering factors used in PyXRD for this study have been set to

280 match this.

281 The kaolinite, illite, talc and chlorite phases are composed of a single component. As such,
282 these are testing the basic aspects of the model such as the orientation factor σ^* , the
283 calculation of the coherent scattering domain size and the calculation of the atomic
284 scattering and structure factors. To test whether PyXRD can handle different sample states
285 correctly, an R0 two-component smectite in air-dried and glycolated state is modelled as
286 well. To further test the implementation of the matrix algorithm for mixed-layer phases, and
287 the related probability models, a number of illite/smectite phases were used. In total seven
288 phases were tested, four of which are two-component illite/smectite phases with
289 Reichweite values varying from 0 to 3 and three of which are three-component
290 illite/smectite phases with Reichweite values varying from 0 to 2. The different smectite
291 components have different hydration states, i.e. the first component always has 2 layers of
292 water (AD state) or 2 layers of glycol molecules (EG state) in its interlayer space while the
293 second component has only a single layer of water or glycol molecules. For these
294 illite/smectites two variants were calculated: one with a low CSDS not at maximum
295 possible degree of ordering (MPDO) and one with a higher CSDS at MPDO.

296 Table 1 contains the Rp factor obtained for these test cases. A few of these patterns are
297 presented in figures 3 and 4. From them and from the Rp and Rwp factors, it is clear
298 PyXRD can produce patterns almost identical to those produced by Sybilla. The small
299 deviations can probably be explained by different physical constants (e.g. atomic
300 scattering factors), although it is impossible to know exactly.

301 **3.2 Comparison between Sybilla and PyXRD results: calculated 00l** 302 **reflections for mixtures of discrete and mixed-layer phyllosilicates**

303 To further validate the model, five patterns were produced in PyXRD for mixtures of

304 increasing complexity. These patterns were imported in Sybilla and modelled using the
305 same phases and the same parameters. This should allow to validate whether the weight
306 fractions in PyXRD can also be obtained by Sybilla. The entered and obtained weight
307 fractions and the corresponding Rp and Rwp factors are presented in table 2. Figure 5
308 shows the comparison between the calculated patterns for mixture 5 from Sybilla and
309 PyXRD. The used phases are largely identical to the phases used in the previous
310 validation, except for the addition of a few phases for which details are also given in table
311 2. The input files for PyXRD and Sybilla are included in this paper's supplement.

312 As can be observed, the weight fractions in PyXRD and Sybilla are reasonably close to
313 each other, with all of the deviations being smaller than 0.5 wt%. These differences can
314 probably, as in the previous validation, be explained by small differences in physical
315 constants used and - in this specific case - unit cell dimensions as well.

316 **3.3 Multispecimen tests**

317 **3.3.1 Assemblage setup**

318 In total, four theoretical mineral assemblages were tested (table 3):

319 Assemblage 1 is a very simple test because of the absence of overlapping and similar
320 phases. Its main purpose was to see whether the program and, more importantly, the
321 selected refinement strategy, can produce a reliable result. The assemblage consists of
322 equal amounts of a discrete kaolinite, a discrete illite and an R0 illite/smectite with only
323 10% illite layers

324 Assemblage 2 is more complex, comprising six different phases: a discrete illite, a
325 discrete kaolinite, an R0 illite/smectite with 65% illite layers, an R0 kaolinite/smectite with
326 80% kaolinite layers, a smectite and a poorly-crystalline chlorite. The idea behind this
327 assemblage was to mimic phases encountered in some soils. The poorly-crystalline

328 chlorite component can be interpreted as a small amount of hydroxy-interlayered smectite
329 (or vermiculite) and is not to be considered a primary trioctahedral chlorite, while the
330 kaolinite/smectite represents a neoformed, defective kaolinite or smectite. This kind of
331 phase has been reported a number of times, usually in finer clay fractions ($\leq 0.2 \mu\text{m}$) of
332 certain soils (Hubert et al., 2009, 2012; Ryan and Huertas, 2009; Dumon et al., 2014). The
333 different phases are also present in different quantities, with the illite-bearing phases each
334 contributing 25.0 wt%, the smectite taking up 20.0 wt%, the kaolinite phases each
335 accounting for 12.5 wt% and the chlorite being a minor phase with only 5.0 wt%.

336 Assemblage 3 is composed of 30% discrete illite, 35% kaolinite, 20% high-charge
337 smectite (vermiculite-like) and 15% low-charge smectite. The main idea behind this test
338 assemblage was to see whether the presence of high-charge and low-charge phases
339 (which in this case produced similar patterns under AD and heated conditions, but different
340 patterns under EG conditions) has an influence on the refinement and the quantification in
341 the different set-ups.

342 Test patterns for assemblage 4 were calculated with 35% well-crystallized kaolinite (with
343 a high average CSDS), 15% poorly-crystallized kaolinite (with a low average CSDS) and
344 50% of an R0 illite/smectite with 98% of illite layers. However, these patterns were not
345 modelled with the same structural models. Instead of two different kaolinites, a single
346 kaolinite was added, and instead of an illite/smectite, a discrete illite was used. As such,
347 the influence of a simplified model input could be checked, which is a common error in
348 real-life uses (e.g. due to misinterpretation).

349 After the necessary phases and their parameters were set up, a calculated pattern was
350 generated from 2 to 50 $^{\circ}2\theta$ with a 0.02° step size, saved and re-imported as experimental
351 data. Random noise was also added to these patterns, using the following formula:

352
$$I_n = I_o \cdot (1 + (X - 0.5) \cdot f_n)$$

353 where I_n is the intensity with noise, I_o the original intensity, X a random fraction between 0
354 and 1 and f_n the noise factor, which was set to 0.01. This results in a random deviation of
355 at most 0.5% above or below the original intensity. This is a high noise level when only
356 considering statistical counting noise, however, these noise levels can be obtained on iron-
357 rich samples when working with a Cu X-ray source due to iron fluorescence. Energy
358 dispersive detectors can eliminate most of this noise nowadays, but it can still be a
359 problem on older equipment, hence it is included here.

360 For assemblages 1 and 2, both the smooth and noisy patterns were used in separate
361 refinements to assess the influence of this treatment. For assemblages 3 and 4, only the
362 noisy patterns were used, because the previous two experiments showed little influence of
363 the noise on the final results (see below).

364 Since evolutionary refinement strategies have a stochastic component, each refinement
365 will be different, even if starting and boundary conditions are identical. Nonetheless, the
366 starting point may also have an influence on the final result. To average out these
367 differences and to check if the final output is reproducible, 50 random starting points were
368 sampled so that a normal distribution over the parameter space was obtained. For each of
369 these points a refinement was started. At the end of these refinements, average parameter
370 values and their standard deviations were calculated for these 50 iterations. Additionally,
371 the model kept track of the best solution found at each generation in each iteration,
372 allowing us to create parameter evolution plots.

373

374 **3.3.2 Assemblage 1**

375 An overview of the obtained average parameter values and standard deviations for
376 assemblage 1 can be found in tables 4 and 5. Parameter evolution plots for two selected
377 parameters (the average CSDS and the fraction of illite layers in the illite/smectite) are also

378 shown in figure 6. Most parameters are determined accurately and with very high
379 precision. The difference between noisy patterns and smooth patterns is marginal, and no
380 difference can be observed between the runs where multiple specimens are combined and
381 those where only a single specimen was used for refinement. As a result of this, the
382 obtained weight fractions for the three phases are also very accurate. The obtained level
383 of accuracy is not a realistic level for natural samples, but stems from the simplicity of this
384 set-up. For the runs using the noisy patterns, a very small (and systematic) deviation in the
385 obtained weight fractions can be observed. This is probably the result of the added noise,
386 since the deviation is not present for runs using the smooth patterns.

387

388 **3.3.3 Assemblage 2**

389 An overview of the obtained average parameter values and standard deviations for
390 assemblage 2 can be found in tables 6 and 7. As was the case in the previous
391 assemblage, no significant difference can be observed between runs that use smooth
392 patterns and those that use noisy ones. Both types produced similar parameter accuracies
393 and precisions. Overall, the results are less accurate and precise compared to
394 assemblage 1, but still very good. Most notably, the weight fractions of the smectite layer
395 types in the kaolinite/smectite show a much larger imprecision. This is also the case in the
396 parameter evolution plots (figure 7) for these fractions. An explanation can be found in the
397 sensitivity of these parameters: since the kaolinite fraction in this mixed-layer is relatively
398 high (80%), the relative amounts of the different types of smectite layers do not have such
399 a large influence on the calculated pattern. Some differences are also noticeable between
400 runs that combine multiple specimens and those where only heated patterns were used.
401 For the latter, the imprecision on the weight fractions for the illite, illite/smectite and
402 smectite phases is significantly larger compared to the other runs. This is to be expected,

403 as heating collapses swelling layers, causing significant peak overlap with the illite peaks.
404 Despite this overlap, it was still possible to obtain accurate and precise averages for the
405 other parameters, comparable to the other runs.

406 **3.3.4 Assemblage 3**

407 An overview of the obtained average parameter values and standard deviations for
408 assemblage 3 can be found in table 8. With this assemblage, the combined set-up and the
409 set-up using only the EG pattern both resulted in the same performance, giving accurate
410 and precise parameter values. The set-up with AD or heated patterns, on the other hand,
411 led to inaccurate and imprecise results, especially when the weight fractions are taken into
412 account. Finally, it can also be observed that the weight fractions and parameter values of
413 phases that were unaffected by the treatments (i.e. kaolinite and illite) are more accurate
414 and precise in these set-ups. It is mainly for the overlapping phases (i.e. smectites) that
415 the errors occur.

416 Figure 8 shows the parameter plots for the multispecimen set-up and the AD set-up for a
417 few selected parameters. This figure illustrates the divergent nature of some parameters in
418 the AD set-up very well, while it is clear that the combined set-up does not suffer from this
419 as it has access to the EG pattern as well.

420 The outcome of this experiment is in line with our expectations, as only the EG pattern
421 contains enough information to distinguish these two smectites from each other. When the
422 EG pattern is absent, the results become divergent, resulting in the high imprecision
423 observed for the AD and heated pattern set-ups.

424 **3.3.5 Assemblage 4**

425 An overview of the obtained average parameter values and standard deviations for
426 assemblage 4 can be found in table 9. In this set-up, we intentionally misidentified a

427 mixed-layer illite/smectite as an illite and overlooked the presence of two populations of
428 kaolinite instead of one. Nevertheless the flawed structural model is able to give us decent
429 parameter accuracies. These kinds of 'mistakes' are quite common in the real-life use of
430 this kind of program, and apparently do not matter too much either, as long as they are
431 related to natural inhomogeneities. In contrast, a model based on a completely wrong
432 interpretation will never yield any good output, and will result in a very obvious mismatch
433 between the calculated and observed patterns. Even in this assemblage, the (residual)
434 XRD patterns (figure 9) show a clear mismatch for these phases. An observant user
435 should notice this and as such be able to identify wrong and/or missing phases.

436 **3.3.6 Summary**

437 For all four assemblages, PyXRD has been able to reproduce the input parameters or at
438 least approximate them with the multispecimen approach. The only complications occur
439 when single patterns are used which do not contain enough information on their own (in
440 most cases heated patterns).

441 The results for these theoretical assemblages seem to suggest that the multi-specimen
442 approach does not add a lot of constraints to the mathematical model. Instead, it appears
443 far more important to correctly identify the phases using multiple specimens than to use
444 these for the parameter refinement. As a result, once the phases are correctly identified, a
445 good quantification can often be obtained with only a single pattern if all phases can be
446 sufficiently discerned from one another in that state. For most natural samples, this could
447 imply that it is sufficient to model the EG and/or the AD pattern. Indeed, many papers
448 presenting modelled X-ray diffraction patterns of phyllosilicates only use the AD and/or EG
449 patterns (Plançon and Roux, 2010; Hubert et al., 2012; Ufer et al., 2012a; Dumon et al.,
450 2014). However, it is important to realize that these results from theoretical experiments
451 cannot be extrapolated automatically to all real-life modelling experiments.

452 In this context, one needs to understand how realistic it is to share some of the parameters
453 between the different specimens during the refinement. Some of them are rather difficult or
454 impossible to control from measurement to measurement. For example, the number of
455 water or glycol layers intercalated into smectite bearing phases is not only dependent on
456 layer charge and the saturating cation, but also on the ambient conditions (i.e. temperature
457 and relative humidity) (Tamura et al., 2000). Because of this, a lot of the parameters
458 cannot and should not be shared, and the advantage of having added more observations
459 is partially lost.

460 **5. Conclusion**

461 In this paper we have presented PyXRD, a new, free and open-source program to perform
462 a (semi-)quantitative analysis of disordered layered minerals using multispecimen X-ray
463 diffraction profile fitting. It is the authors' sincere hope that others will pick up on the
464 program and improve it. The novelty of this program lies specifically in the *ab initio*
465 incorporation of the multispecimen method, making it possible to share phases and (a
466 selection of) their parameters across multiple specimens. This allows to model several
467 specimens side-by-side, and is an important step forward. In theory, this could also help in
468 further constraining the mathematical model and thus improving the automatic parameter
469 refinement results (Sakharov et al., 1999a; Meunier, 2005; Lanson, 2011). However,
470 results from theoretical experiments indicate that a multispecimen refinement setup is not
471 always required to obtain good parameter estimates. Rather, it is far more important to use
472 the multispecimen method to obtain a correct identification of the phases present. We can
473 conclude that PyXRD has proven to be very useful when X-ray diffraction patterns for
474 complex mineral assemblages containing (mixed-layer) phyllosilicates are modelled with a
475 multispecimen approach.

476 **6. Code availability**

477 The source code for PyXRD can be found online at [https://github.com/mathijs-](https://github.com/mathijs-dumon/PyXRD)
478 [dumon/PyXRD](https://github.com/mathijs-dumon/PyXRD), together with installation instructions and a manual with detailed
479 information regarding the calculations and a step-by-step example on how to use the user
480 interface.

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Table 1: Overview of the discrete phases used to compare the output from PyXRD with the output of Sybilla (R is Reichweite, G is the number of components, \bar{N} is the average CSDS, AD is air-dry, EG is ethylene-glycol, relevant probability (P) and weight (W) factors are given, Rp and Rwp are the unweighted and weighted residual errors of the patterns respectively).

| | Phase | R | G | State | \bar{N} | P and W factors | | | Rp | Rwp |
|-----|-----------------|---|---|-------|-----------|---|--|--|-----|-----|
| 1 | Kaolinite | - | 1 | - | 20 | - | | | 0.7 | 0.9 |
| 2 | Illite | - | 1 | - | 20 | - | | | 0.9 | 1.3 |
| 3 | Talc | - | 1 | - | 20 | - | | | 0.8 | 1.0 |
| 4 | Chlorite | - | 1 | - | 20 | - | | | 0.8 | 1.0 |
| 5 | Illite/Smectite | 0 | 2 | AD | 4 | $W_1 = 0.5$ | | | 1.0 | 1.6 |
| 6 | Illite/Smectite | 1 | 2 | AD | 4 | $W_1 = 0.6$ $P_{22} = 0.5$ | | | 1.0 | 1.6 |
| 6b | Illite/Smectite | 1 | 2 | AD | 15 | $W_1 = 0.6$ $P_{22} = 0.0$ | | | 0.7 | 1.5 |
| 7 | Illite/Smectite | 2 | 2 | AD | 4 | $W_1 = 0.6$ $P_{21} = 1.0$ | $P_{112} = 0.5$ $P_{221} = 0.0$ | | 1.4 | 2.1 |
| 7b | Illite/Smectite | 2 | 2 | AD | 15 | $W_1 = 0.6$ $P_{21} = 1.0$ | $P_{112} = 1.0$ $P_{221} = 0.0$ | | 0.7 | 1.5 |
| 8 | Illite/Smectite | 3 | 2 | AD | 4 | $W_1 = 0.9$ $P_{2112} = 0.5$ | $P_{22}=0$ $P_{212}=0$ | | 1.9 | 2.3 |
| 8b | Illite/Smectite | 3 | 2 | AD | 15 | $W_1 = 0.9$ $P_{2112} = 0.0$ | $P_{22}=0$ $P_{212}=0$ | | 0.5 | 1.0 |
| 9 | Illite/Smectite | 0 | 3 | AD | 4 | $W_1 = 0.33$ $W_2 / (W_2 + W_3) = 0.5$ | | | 1.1 | 1.7 |
| 10 | Illite/Smectite | 1 | 3 | AD | 4 | $W_1 = 0.5$ $W_2 / (W_2 + W_3) = 0.8$ | $P_{11} = 0.85$ $(W_{22} + W_{23}) / (W_{22} + W_{23} + W_{32} + W_{33}) = 0.85$ | $W_{22} / (W_{22} + W_{23}) = 0.8$ $W_{32} / (W_{32} + W_{33}) = 0.7$ | 1.1 | 1.7 |
| 10b | Illite/Smectite | 1 | 3 | AD | 15 | $W_1 = 0.5$ $W_2 / (W_2 + W_3) = 0.8$ | $P_{11} = 0.0$ $(W_{22} + W_{23}) / (W_{22} + W_{23} + W_{32} + W_{33}) = 0.85$ | $W_{22} / (W_{22} + W_{23}) = 0.8$ $W_{32} / (W_{32} + W_{33}) = 0.7$ | 1.0 | 1.7 |
| 11 | Illite/Smectite | 2 | 3 | AD | 4 | $W_1 = 0.8$ $W_2 / (W_2 + W_3) = 0.5$ | $P_{x1x} = 0.5$ $(W_{212} + W_{213}) / (W_{212} + W_{213} + W_{312} + W_{313}) = 0.5$ | $W_{212} / (W_{212} + W_{213}) = 0.5$ $W_{312} / (W_{312} + W_{313}) = 0.5$ | 1.7 | 2.2 |
| 11b | Illite/Smectite | 2 | 3 | AD | 15 | $W_1 = 0.8$ $W_2 / (W_2 + W_3) = 0.5$ | $P_{x1x} = 0.0$ $(W_{212} + W_{213}) / (W_{212} + W_{213} + W_{312} + W_{313}) = 0.5$ | $W_{212} / (W_{212} + W_{213}) = 0.5$ $W_{312} / (W_{312} + W_{313}) = 0.5$ | 0.5 | 1.0 |
| 12 | Smectite | 0 | 2 | AD | 4 | $W_1 = 0.7$ | | | 1.3 | 1.7 |
| 13 | Smectite | 0 | 2 | EG | 4 | $W_1 = 0.7$ | | | 1.0 | 1.2 |

Table 2: Overview of the test mixtures used to compare the weight fraction output from PyXRD with the output of Sybilla, with details for the different phases (R is Reichweite, \bar{N} is the average CSDS, d_{001} is the basal spacing, relevant probability (P) and weight (W) factors are given).

| Mixture | Rp | Rwp | Phases | PyXRD wt% | Sybilla wt% | Phase characteristics |
|---------|-----|-----|-------------|--------------|----------------|---|
| 1 | 0.8 | 1.0 | Kaolinite | 70.0 | 69.9 | As in table 1 |
| | | | Illite | 30.0 | 30.1 | As in table 1 |
| 2 | 1.1 | 1.5 | Kaolinite | 20.0 | 20.0 | As in table 1 |
| | | | Illite | 30.0 | 30.1 | As in table 1 |
| | | | IS R0 | 10.0 | 9.5 | As in table 1 |
| | | | SSS R0 | 40.0 | 40.5 | $\bar{N} = 4$; $W_1=0.8$; $W_2/(W_2+W_3)=0.8$ |
| 3 | 1.0 | 1.4 | Kaolinite | 10.0 | 10.0 | As in table 1 |
| | | | Illite | 25.0 | 25.1 | As in table 1 |
| | | | Chlorite | 20.0 | 20.1 | As in table 1 |
| | | | IS R0 | 15.0 | 14.7 | As in table 1 |
| | | | CS R1 | 10.0 | 10.1 | $\bar{N} = 10$; $W_1=0.5$; $P_{11} = 0.1$ |
| | | | SSS R0 | 20.0 | 20.0 | $\bar{N} = 4$; $W_1=0.8$; $W_2/(W_2+W_3)=0.8$ |
| 4 | 1.3 | 1.8 | ISS R0 | 15.0 | 15.2 | As in table 1 |
| | | | CSS R0 | 5.0 | 5.0 | $\bar{N} = 5$; $W_1=0.4$; $W_2/(W_2+W_3)=0.9$ |
| | | | Chlorite | 5.0 | 5.0 | As in table 1 |
| | | | Illite | 15.0 | 14.9 | As in table 1 |
| | | | Kaolinite 1 | 15.0 | 14.9 | As in table 1 |
| | | | Kaolinite 2 | 25.0 | 25.0 | $\bar{N} = 6$; $d_{001} = 0.718$ nm |
| | | | IS R1 | 10.0 | 10.0 | As in table 1 |
| | | | CS R1 | 10.0 | 10.1 | $\bar{N} = 10$; $W_1=0.5$; $P_{11} = 0.1$ |
| 5 | 1.7 | 2.4 | ISS R0 | 10.0 | 10.0 | As in table 1 |
| | | | CSS R0 | 10.0 | 10.0 | $\bar{N} = 5$; $W_1=0.4$; $W_2/(W_2+W_3)=0.9$ |
| | | | Chlorite | 10.0 | 10.0 | As in table 1 |
| | | | Illite | 10.0 | 9.9 | As in table 1 |
| | | | Kaolinite 1 | 10.0 | 10.0 | As in table 1 |
| | | | Kaolinite 2 | 10.0 | 10.0 | $\bar{N} = 6$; $d_{001} = 0.718$ nm |
| | | | IS R1 | 10.0 | 10.1 | As in table 1 |
| | | | CS R1 | 10.0 | 9.9 | $\bar{N} = 10$; $W_1=0.5$; $P_{11} = 0.1$ |
| | | | SS R0 | 10.0 | 9.9 | $\bar{N} = 4$; $W_1=0.7$ |
| | | | KSS R0 | 10.0 | 10.0 | $\bar{N} = 7$; $W_1=0.6$; $W_2/(W_2+W_3)=0.8$ |

Table 3: Overview of the different test assemblages for the comparison between multispecimen and single pattern refinements and the type of refined patterns.

| | Assemblage | Smooth pattern? | Noisy pattern? |
|---|---|------------------------|-----------------------|
| 1 | 33.3% Kaolinite 33.3% Illite 33.3% Illite/Smectite (10/90) R0 | yes | yes |
| 2 | 25.0% Illite 25.0% Illite/Smectite (65/35) R0 20.0% Smectite 12.5% Kaolinite 12.5% Kaolinite/Smectite (80/20) R0 5.0% Chlorite | yes | yes |
| 3 | 35.0% Kaolinite 30.0% Illite 15.0% High-charge smectite 20.0% Low-charge smectite | no | yes |
| 4 | 35.0% Kaolinite (CSDS = 20) 15.0% Kaolinite (CSDS = 6) 50.0% Illite/Smectite (98/2) R0 | no | yes |

Table 4: Overview of the means and standard deviations for weight fractions and refined parameters for assemblage 1 using smooth patterns.

| Assemblage #1 - smooth patterns | | | Range | | Multiple specimens (n=50) | Only AD (n=50) | Only EG (n=50) | Only 350 heated (n=50) |
|---------------------------------|-----------------------|------------|-------|------|------------------------------------|------------------------------------|------------------------------------|------------------------------------|
| Phase | Property name | True value | Min. | Max. | Obtained value $\mu \pm \sigma$ | Obtained value $\mu \pm \sigma$ | Obtained value $\mu \pm \sigma$ | Obtained value $\mu \pm \sigma$ |
| Kaolinite | wt% | 33.3 | – | – | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 |
| | T | 10.0 | 8.0 | 20.0 | 10.0 ± 0.00 | 10.0 ± 0.00 | 10.0 ± 0.00 | 10.0 ± 0.00 |
| Illite | wt% | 33.3 | – | – | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 |
| | T | 10.0 | 8.0 | 20.0 | 10.0 ± 0.00 | 10.0 ± 0.00 | 10.0 ± 0.00 | 10.0 ± 0.00 |
| Illite/Smectite R0 | wt% | 33.3 | – | – | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 | 33.3 ± 0.00 |
| | T | 5.0 | 3.0 | 10.0 | 5.0 ± 0.00 | 5.0 ± 0.00 | 5.0 ± 0.00 | 5.0 ± 0.00 |
| | Illite content | 0.1 | 0.0 | 1.0 | 0.10 ± 0.00 | 0.10 ± 0.00 | 0.10 ± 0.00 | 0.10 ± 0.00 |
| | 2wat / (2wat + 1wat) | 0.5 | 0.0 | 1.0 | 0.50 ± 0.00 | 0.50 ± 0.00 | – | – |
| | 2 alv / (2alv + 1alv) | 0.5 | 0.0 | 1.0 | 0.50 ± 0.00 | – | 0.50 ± 0.00 | – |
| | 0 alv / (0alv + 1alv) | 1.0 | 0.0 | 1.0 | 1.00 ± 0.00 | – | – | 1.00 ± 0.00 |

Table 6: Overview of the means and standard deviations of weight fractions and refined parameters for assemblage 2 using smooth patterns.

| <i>Assemblage #2 – Smooth patterns</i> | | | | | Multiple specimens (n=50) | Only AD (n=50) | Only EG (n=50) | Only 350°C (n=50) |
|--|-------------------------------|------------|-------|------|------------------------------|-------------------|-------------------|----------------------|
| Phase | Property name | True value | Range | | Obtained value | Obtained value | Obtained value | Obtained value |
| | | | Min. | Max. | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ |
| Illite | wt% | 25.0 | – | – | 25.0 ± 0.1 | 25.0 ± 0.1 | 25.0 ± 0.0 | 25.4 ± 0.71 |
| | T | 13.0 | 10.0 | 30.0 | 13.0 ± 0.1 | 13.0 ± 0.0 | 13.0 ± 0.0 | 12.9 ± 0.2 |
| Illite/Smectite R0 | wt% | 25.0 | – | – | 24.9 ± 0.2 | 25.0 ± 0.1 | 25.0 ± 0.0 | 24.8 ± 0.3 |
| | T | 5.0 | 3.0 | 10.0 | 5.1 ± 0.1 | 5.0 ± 0.0 | 5.0 ± 0.0 | 5.0 ± 0.1 |
| | Illite content | 0.65 | 0.5 | 1.0 | 0.65 ± 0.00 | 0.65 ± 0.00 | 0.65 ± 0.00 | 0.64 ± 0.03 |
| | 2wat / (2wat + 1wat) | 0.7 | 0.0 | 1.0 | 0.70 ± 0.01 | 0.70 ± 0.00 | – | – |
| | 2 alv / (2alv + 1alv) | 0.7 | 0.0 | 1.0 | 0.71 ± 0.02 | – | 0.70 ± 0.00 | – |
| | 0 alv / (0alv + 1alv) | 1.0 | 0.8 | 1.0 | 0.96 ± 0.03 | – | – | 0.99 ± 0.01 |
| Kaolinite | wt% | 12.5 | – | – | 12.5 ± 0.0 | 12.5 ± 0.0 | 12.5 ± 0.0 | 12.5 ± 0.0 |
| | T | 20.0 | 10.0 | 30.0 | 20.1 ± 0.1 | 20.0 ± 0.0 | 20.0 ± 0.0 | 20.1 ± 0.1 |
| Kaolinite/Smectite R0 | wt% | 12.5 | – | – | 12.7 ± 0.2 | 12.5 ± 0.1 | 12.5 ± 0.0 | 12.9 ± 0.2 |
| | T | 3.0 | 3.0 | 10.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 |
| | Kaolinite content | 0.80 | 0.7 | 1.0 | 0.80 ± 0.01 | 0.80 ± 0.00 | 0.80 ± 0.00 | 0.79 ± 0.00 |
| | 2wat / (2wat + 1wat) | 0.25 | 0.0 | 0.6 | 0.26 ± 0.11 | 0.25 ± 0.02 | – | – |
| | 2 alv / (2alv + 1alv) | 0.50 | 0.0 | 0.6 | 0.44 ± 0.10 | – | 0.50 ± 0.01 | – |
| | 0 alv / (0alv + 1alv) | 1.00 | 0.8 | 1.0 | 0.93 ± 0.05 | – | – | 0.93 ± 0.04 |
| Smectite | wt% | 20.0 | – | – | 19.9 ± 0.1 | 20.0 ± 0.1 | 20.0 ± 0.0 | 19.6 ± 0.7 |
| | T | 3.0 | 3.0 | 10.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 |
| | 2wat / (2wat + 1wat) | 0.60 | 0.5 | 1.0 | 0.60 ± 0.00 | 0.60 ± 0.00 | – | – |
| | 2 alv / (2alv + 1alv) | 0.90 | 0.5 | 1.0 | 0.90 ± 0.00 | – | 0.90 ± 0.00 | – |
| | 0 alv / (0alv + 1alv) | 0.90 | 0.8 | 1.0 | 0.92 ± 0.01 | – | – | 0.90 ± 0.01 |
| Chlorite | wt% | 5.0 | – | – | 5.0 ± 0.0 | 5.0 ± 0.1 | 5.0 ± 0.0 | 5.0 ± 0.0 |
| | T | 5.0 | 3 | 10 | 5.0 ± 0.0 | 5.0 ± 0.0 | 5.0 ± 0.0 | 5.0 ± 0.0 |
| | $\partial d_{001} \cdot 10^3$ | 5.0 | 1.0 | 10.0 | 5.0 ± 0.1 | 5.0 ± 0.1 | 5.0 ± 0.0 | 5.1 ± 0.1 |

Table 7: Overview of the means and standard deviations of weight fractions and refined parameters for assemblage 2 using noisy patterns.

| <i>Assemblage #2 – Noisy patterns</i> | | | | | Multiple specimens (n=50) | Only AD (n=50) | Only EG (n=50) | Only 350°C (n=50) |
|---------------------------------------|-------------------------------|------------|-------|------|------------------------------|-------------------|-------------------|----------------------|
| Phase | Property name | True value | Range | | Obtained value | Obtained value | Obtained value | Obtained value |
| | | | Min. | Max. | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ |
| Illite | wt% | 25.0 | – | – | 25.1 ± 0.2 | 25.2 ± 0.1 | 25.3 ± 0.1 | 24.8 ± 1.5 |
| | T | 13.0 | 10.0 | 30.0 | 13.1 ± 0.1 | 13.2 ± 0.0 | 12.9 ± 0.0 | 13.2 ± 0.3 |
| Illite/Smectite R0 | wt% | 25.0 | – | – | 24.6 ± 0.4 | 25.8 ± 0.2 | 24.7 ± 0.1 | 25.8 ± 1.9 |
| | T | 5.0 | 3.0 | 10.0 | 5.0 ± 0.1 | 5.2 ± 0.0 | 4.9 ± 0.0 | 5.0 ± 0.4 |
| | Illite content | 0.65 | 0.5 | 1.0 | 0.64 ± 0.01 | 0.65 ± 0.00 | 0.65 ± 0.00 | 0.64 ± 0.04 |
| | 2wat / (2wat + 1wat) | 0.7 | 0.0 | 1.0 | 0.67 ± 0.02 | 0.70 ± 0.01 | – | – |
| | 2 alv / (2alv + 1alv) | 0.7 | 0.0 | 1.0 | 0.68 ± 0.01 | – | 0.67 ± 0.00 | – |
| | 0 alv / (0alv + 1alv) | 1.0 | 0.8 | 1.0 | 0.96 ± 0.02 | – | – | 0.96 ± 0.03 |
| Kaolinite | wt% | 12.5 | – | – | 12.5 ± 0.0 | 12.3 ± 0.0 | 12.5 ± 0.0 | 12.6 ± 0.1 |
| | T | 20.0 | 10.0 | 30.0 | 20.1 ± 0.1 | 20.1 ± 0.0 | 20.1 ± 0.0 | 20.0 ± 0.0 |
| Kaolinite/Smectite R0 | wt% | 12.5 | – | – | 12.8 ± 0.4 | 12.1 ± 0.2 | 12.4 ± 0.1 | 12.5 ± 0.1 |
| | T | 3.0 | 3.0 | 10.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 |
| | Kaolinite content | 0.80 | 0.7 | 1.0 | 0.80 ± 0.01 | 0.81 ± 0.01 | 0.81 ± 0.00 | 0.82 ± 0.00 |
| | 2wat / (2wat + 1wat) | 0.25 | 0.0 | 0.6 | 0.30 ± 0.11 | 0.34 ± 0.03 | – | – |
| | 2 alv / (2alv + 1alv) | 0.50 | 0.0 | 0.6 | 0.47 ± 0.10 | – | 0.54 ± 0.02 | – |
| | 0 alv / (0alv + 1alv) | 1.00 | 0.8 | 1.0 | 0.91 ± 0.05 | – | – | 0.94 ± 0.04 |
| Smectite | wt% | 20.0 | – | – | 20.1 ± 0.2 | 19.6 ± 0.2 | 20.2 ± 0.1 | 19.5 ± 3.4 |
| | T | 3.0 | 3.0 | 10.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.0 | 3.0 ± 0.2 |
| | 2wat / (2wat + 1wat) | 0.60 | 0.5 | 1.0 | 0.60 ± 0.01 | 0.60 ± 0.00 | – | – |
| | 2 alv / (2alv + 1alv) | 0.90 | 0.5 | 1.0 | 0.90 ± 0.01 | – | 0.90 ± 0.00 | – |
| | 0 alv / (0alv + 1alv) | 0.90 | 0.8 | 1.0 | 0.92 ± 0.01 | – | – | 0.91 ± 0.02 |
| Chlorite | wt% | 5.0 | – | – | 5.0 ± 0.0 | 5.1 ± 0.1 | 4.9 ± 0.0 | 4.9 ± 0.1 |
| | T | 5.0 | 3 | 10 | 5.1 ± 0.0 | 5.2 ± 0.1 | 5.2 ± 0.0 | 5.0 ± 0.0 |
| | $\partial d_{001} \cdot 10^3$ | 5.0 | 1.0 | 10.0 | 5.2 ± 0.3 | 5.5 ± 0.3 | 4.5 ± 0.2 | 5.4 ± 0.3 |

Table 8: Overview of the means and standard deviations of weight fractions and refined parameters for assemblage 3.

| <i>Assemblage #3 – Noisy patterns</i> | | | | | Multiple specimens (n=50) | Only AD (n=50) | Only EG (n=50) | Only 350°C (n=50) |
|---------------------------------------|----------------|------------|-------|------|------------------------------|-------------------|-------------------|----------------------|
| Phase | Property name | True value | Range | | Obtained value | Obtained value | Obtained value | Obtained value |
| | | | Min. | Max. | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ |
| Kaolinite | wt% | 35.0 | – | – | 35.0 ± 0.0 | 35.3 ± 0.6 | 34.7 ± 0.0 | 34.9 ± 0.0 |
| | T | 18.0 | 5 | 40 | 18.0 ± 0.0 | 18.0 ± 0.2 | 18.0 ± 0.0 | 18.0 ± 0.0 |
| Illite | wt% | 30.0 | – | – | 30.0 ± 0.0 | 30.0 ± 0.8 | 30.1 ± 0.1 | 29.0 ± 0.1 |
| | T | 25.0 | 5 | 40 | 25.0 ± 0.0 | 25.5 ± 0.1 | 24.8 ± 0.0 | 25.2 ± 0.12 |
| High-charge smectite | wt% | 15.0 | – | – | 15.1 ± 0.0 | 16.9 ± 5.9 | 15.8 ± 0.1 | 16.0 ± 0.1 |
| | T | 10.0 | 5 | 40 | 10.0 ± 0.0 | 11.3 ± 5.2 | 10.0 ± 0.0 | 10.0 ± 0.1 |
| | HC / (HC + LC) | 0.90 | 0.50 | 1.00 | 0.90 ± 0.00 | 0.87 ± 0.06 | 0.90 ± 0.00 | – |
| Low-charge smectite | wt % | 20.0 | – | – | 19.9 ± 0.0 | 17.8 ± 7.2 | 19.4 ± 0.2 | 20.3 ± 0.2 |
| | T | 10.0 | 5 | 40 | 10.0 ± 0.0 | 12.2 ± 7.4 | 10.0 ± 0.0 | 10.2 ± 0.1 |
| | LC / (LC + HC) | 0.80 | 0.50 | 1.00 | 0.80 ± 0.00 | 0.83 ± 0.06 | 0.80 ± 0.00 | – |

Table 9: Overview of the means and standard deviations of weight fractions and refined parameters for assemblage 4.

| <i>Assemblage #4 – Noisy patterns</i> | | | | | Multiple specimens (n=50) | Only AD (n=50) | Only EG (n=50) | Only 350°C (n=50) |
|---------------------------------------|---|------------|-------|------|------------------------------|-------------------|-------------------|----------------------|
| Phase | Property name | True value | Range | | Obtained value | Obtained value | Obtained value | Obtained value |
| | | | Min. | Max. | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ | $\mu \pm \sigma$ |
| Kaolinite | wt% | 50.0 | – | – | 49.7 ± 0.1 | 49.3 ± 0.0 | 50.3 ± 0.2 | 49.3 ± 0.1 |
| | T | 15.8 | 5 | 40 | 15.2 ± 0.1 | 15.2 ± 0.0 | 15.2 ± 0.0 | 15.6 ± 0.0 |
| Illite | wt% | 50.0 | – | – | 50.3 ± 0.1 | 50.7 ± 0.0 | 49.7 ± 0.2 | 50.7 ± 0.1 |
| | T | 30.0 | 5 | 40 | 21.2 ± 0.0 | 18.8 ± 0.0 | 22.7 ± 0.1 | 28.0 ± 0.0 |
| | Oct. Fe ³⁺ / Oct. Al ³⁺ | 0.125 | 0 | 0.5 | 0.133 ± 0.000 | 0.126 ± 0.002 | 0.151 ± 0.001 | 0.139 ± 0.001 |
| | K content | 1.50 | 0 | 2 | 1.52 ± 0.01 | 1.49 ± 0.00 | 1.52 ± 0.01 | 1.44 ± 0.00 |

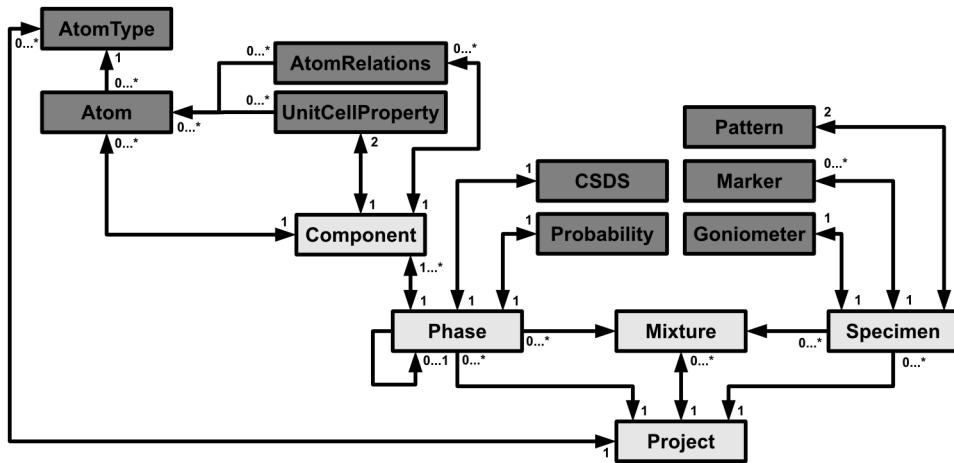


Figure 1: Schematic overview of the most important objects in PyXRD and their relations. Arrows indicate 'is referenced x times by' relations and the numbers indicate the multiplicity of that relation (e.g. Project holds 0 or more references to AtomType).

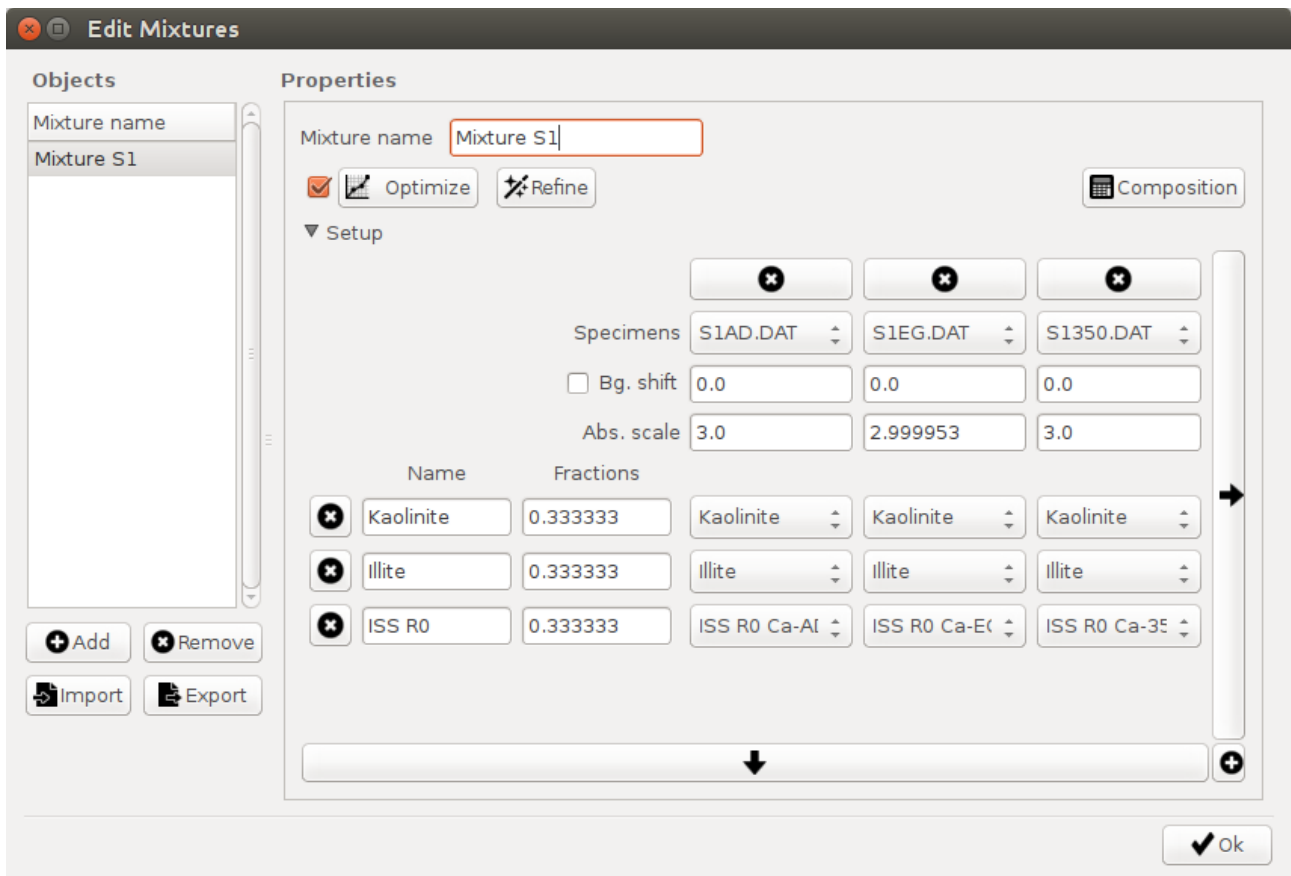


Figure 2: screenshot showing the 'Edit mixtures' dialog where a user can link different phases (Kaolinite, Illite, ISS R0 Ca-AD, ...) with the corresponding specimens (S1AD.dat, S1EG.dat, ...).

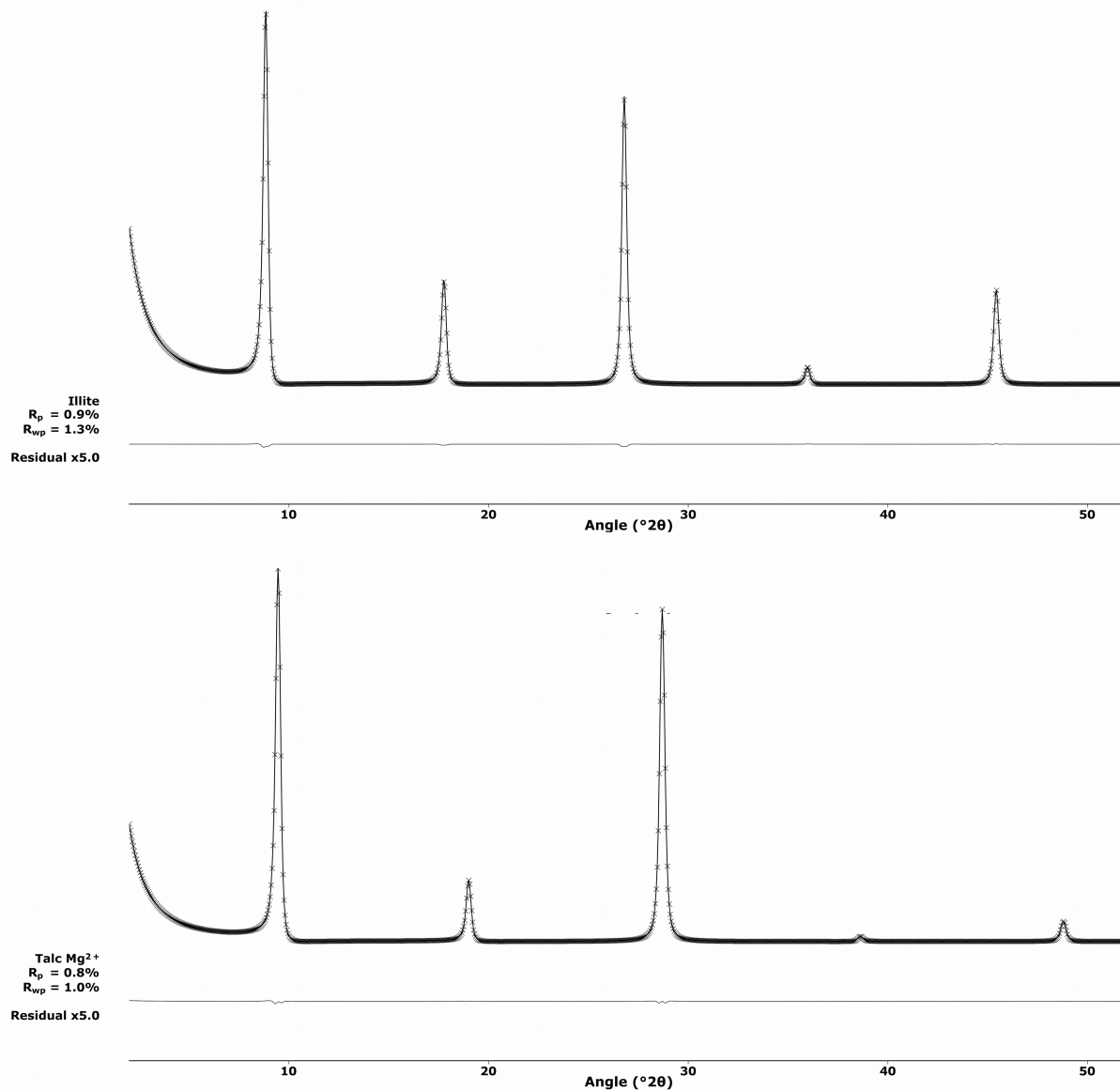


Figure 3 – Calculated patterns for discrete illite (top) and talc (bottom), showing nearly identical output for PyXRD (solid line) and Sybilla (crosses). For clarity the residual patterns are scaled to 5x their original intensity.

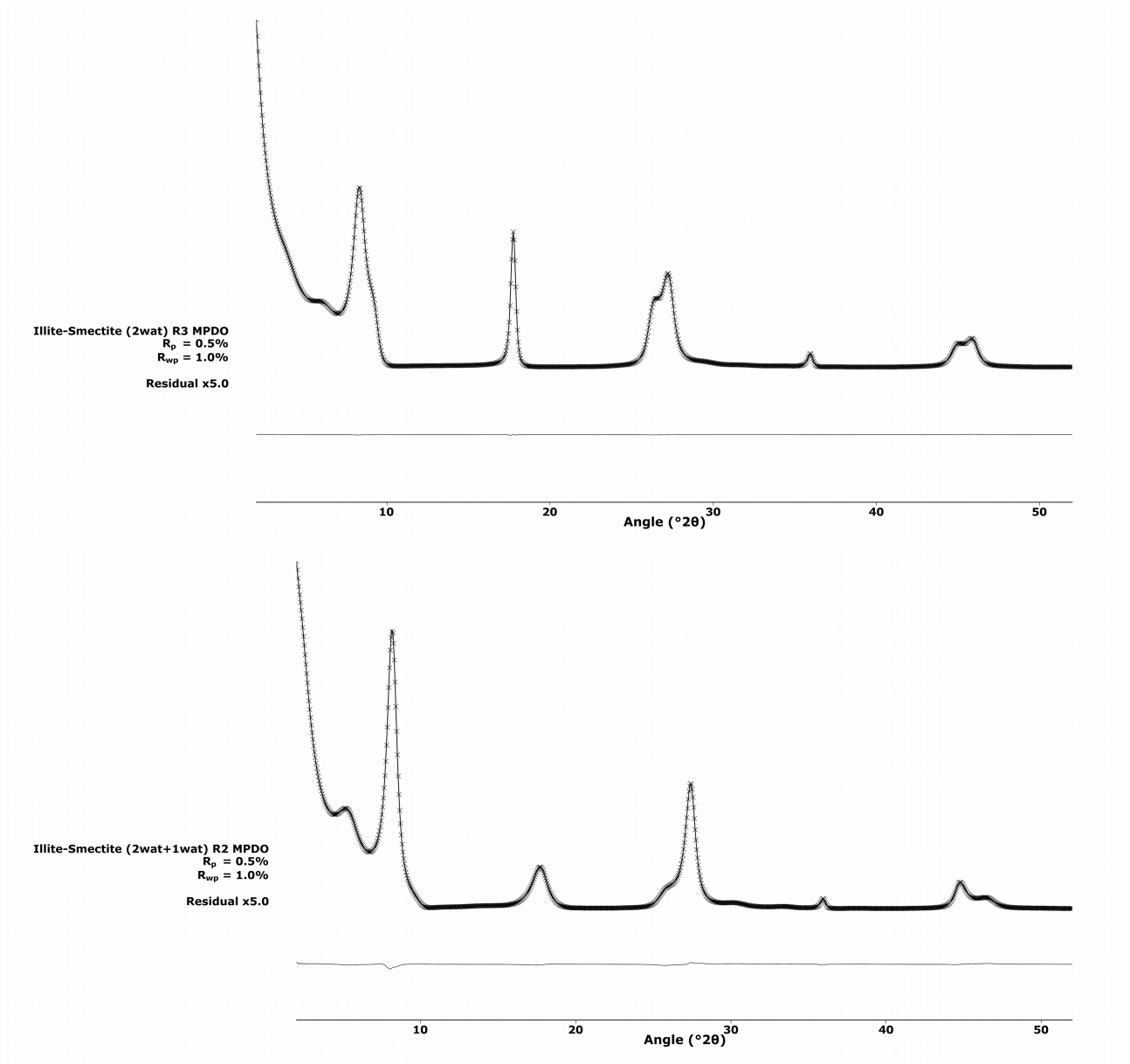


Figure 4 – Calculated patterns for illite-smectite (IS) R3 (top) and illite-smectite (ISS) R2 (bottom; both have MPDO), showing nearly identical output for PyXRD (solid line) and Sybilla (crosses). For clarity the residual patterns are scaled to 5x their original intensity.

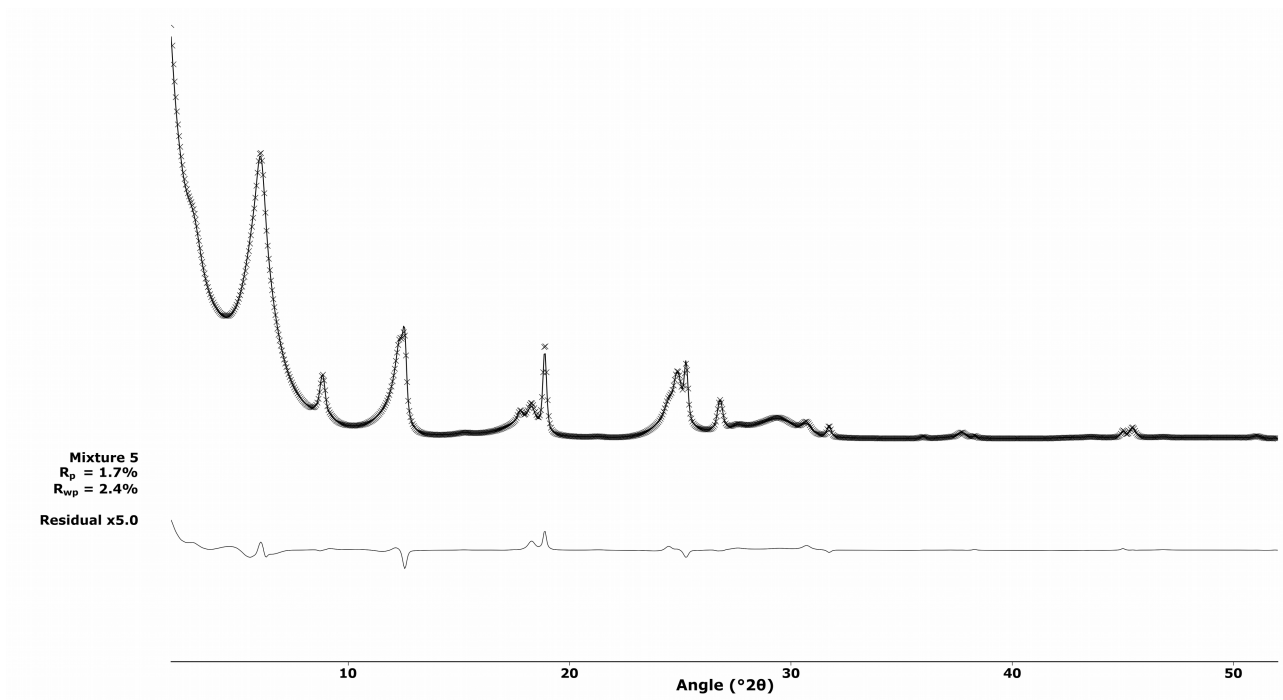


Figure 5 – Calculated patterns for mixture 5 for PyXRD (solid line) and Sybilla (crosses). For clarity the residual pattern is scaled to 5x its original intensity.

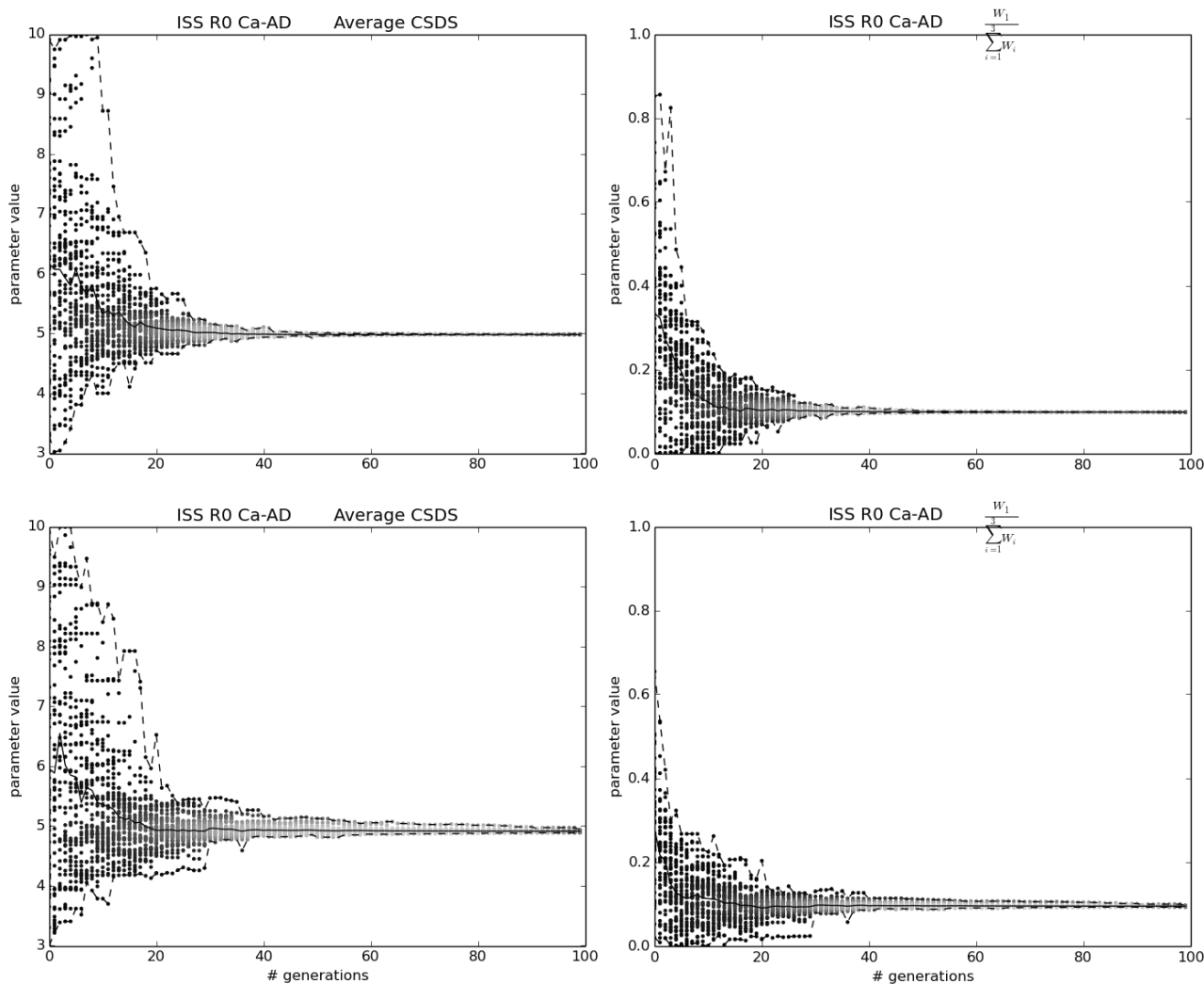


Figure 6: Parameter evolution plots (left: average CSDS; right: illite content) for the noisy patterns of assemblage 1 for the multispecimen run (top plots) and the isolated AD run (bottom plots). Minimum and maximum values during the refinement are indicated with dashed lines, iterations' best solutions at each generation indicated by dots and average solution with a solid line. The higher the density of the dots, the lighter they are colored.

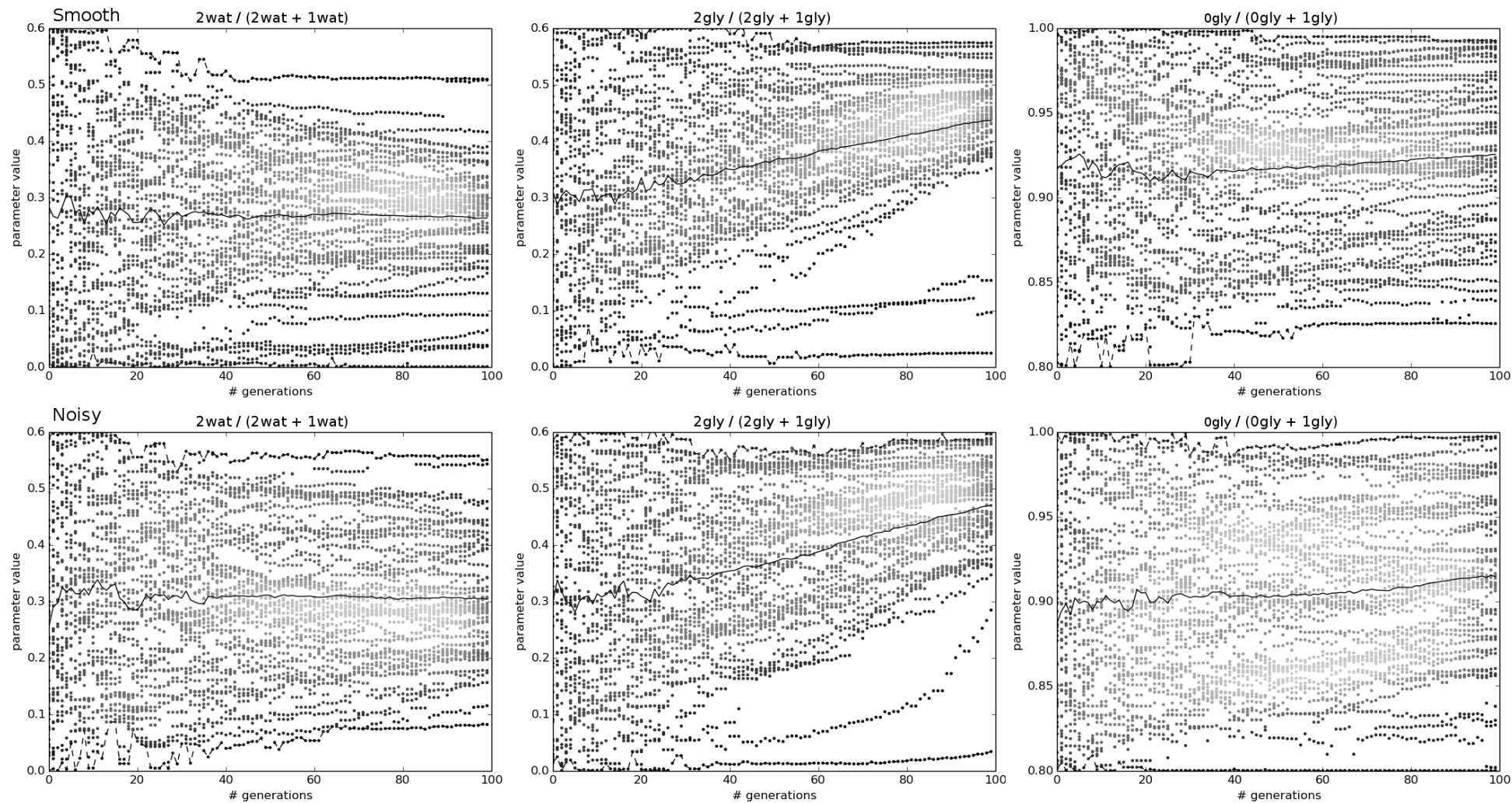


Figure 7: Parameter evolution plots for the smectite fractions in the kaolinite-smectite mixed layer of assemblage 2 using the multispecimen setup. Plots for the smooth patterns are in the top row, for noisy patterns in the bottom row. Legend as in Figure 6.

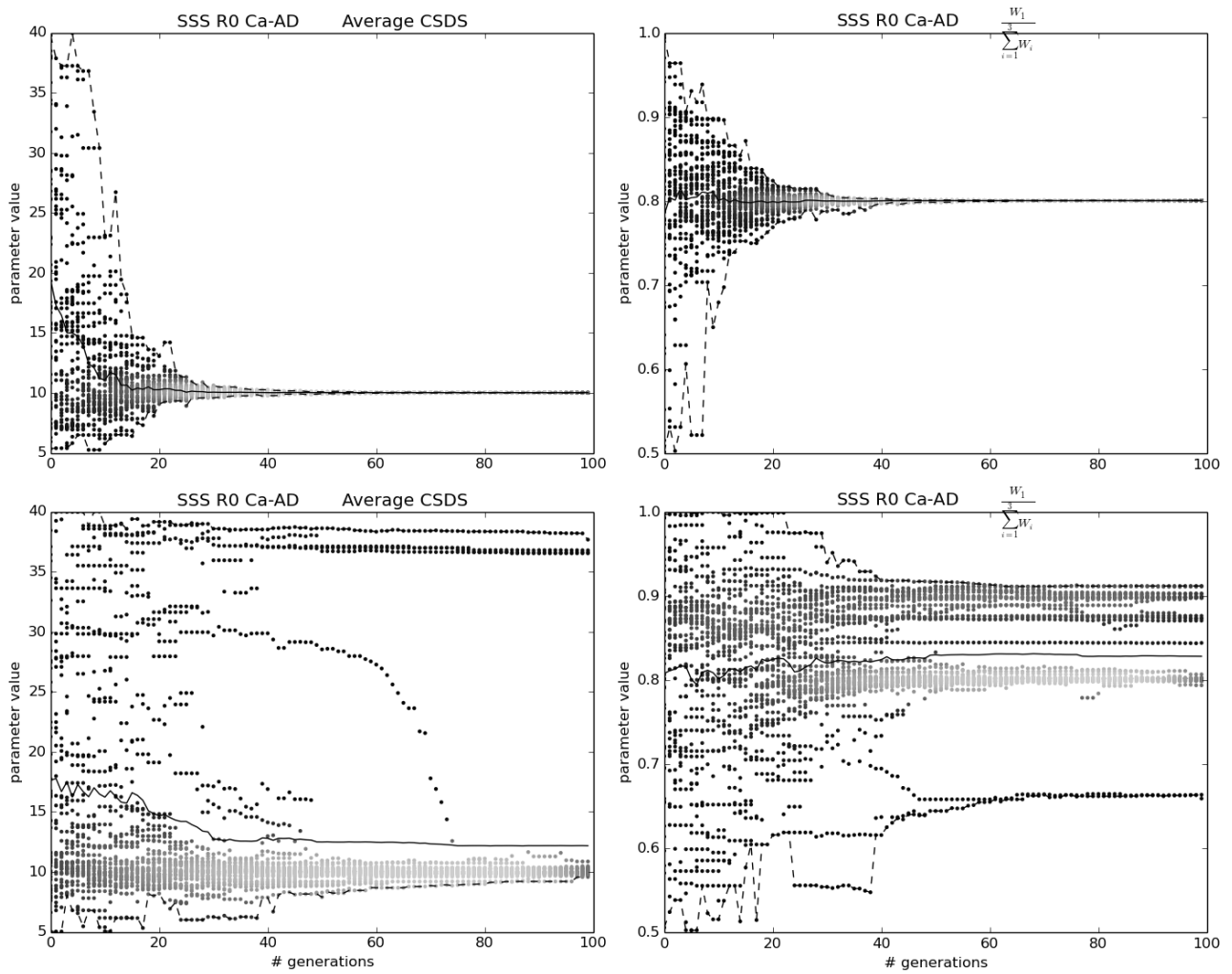


Figure 8: Parameter evolution plots for the low-charge smectite in assemblage 3. Plots for the multispecimen setup are in the top row, for the AD single pattern setup in the bottom row. Legend as in Figure 6.

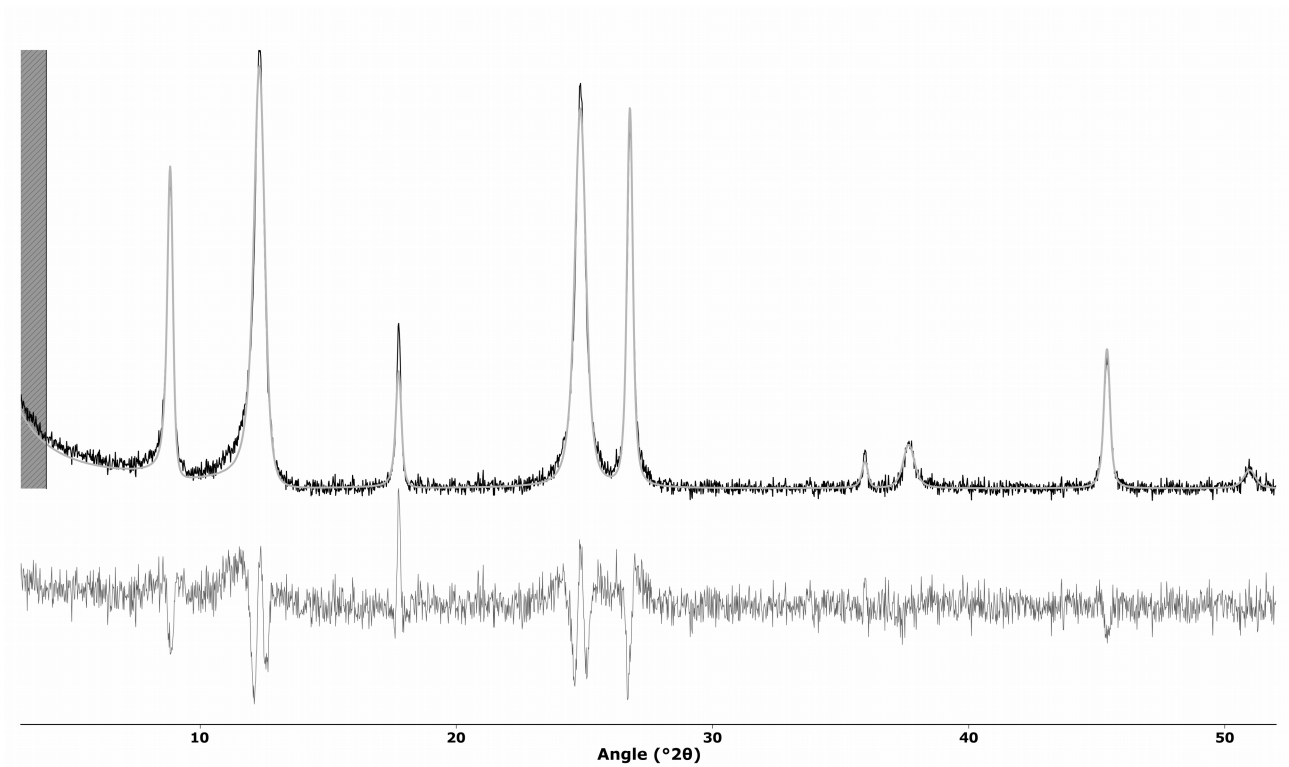


Figure 9: The input (black solid line) and refined (grey solid line) AD pattern and their difference (grey solid line at the bottom) for the multispecimen setup of assemblage 4. An observant user should see the mismatches in the patterns and realize his model needs improvement.