

Supplementary information for the manuscript

“Fast energy minimization of the CCDC drug-subset structures by molecule-in-cluster computations allows independent structure validation and model completion”

by

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Content:

- List of 53 Drug Subset¹ structures with atoms or molecules on special positions not considered in our GFN2-xTB² computations
- List of disordered structures where two separate conformers were optimized
- List of structures where optimization failed or was not performed due to missing atoms combined with structural complexity.
- List of 708 optimized structures. When `_m` is added to the refcode it means that the molecule were either not within the unit cell, or needed moving to bring cation and anion close together.
- List of 690 RMSD values for most of the optimized structures that serve as an additional cross validation and confirm that the optimized structure agrees with the experimental one. These were computed between the optimized and the original refcode coordinates, using the non-hydrogen atom coordinates only. The flagged entries were checked and corrected afterwards. This illustrates the power of combining quantum chemical optimization with experiment.
- Computer-generated list of 732 plots of ASU-in-cluster geometry optimizations, displaying energy versus individual optimization cycle, re-generating the cluster four times each time

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Table 1: List of 53 Drug Subset¹ structures with atoms or molecules on special positions not considered in our GFN2-xTB² computations

1. ACXMAC_water_on_special_position
2. AMALAC_1_molecule_on_special_position
3. ASATOD_mol_on_special_position
4. AWUWIY01_water_on_special_position
5. BETXON_water_on_special_position
6. BINZIH_cl_on_special_position
7. BOKHAM02_biphenyl_on_special_position
8. BRPRBZ_molecule_on_special_position
9. CAASCO02_Ca_on_special_position
10. CATDAE_mol_on_special_position
11. CIGCIG_molecule_on_special_position
12. CYCLIZ10_molecule_on_special_position
13. CYPHEP_water_on_special_position
14. EPUDIE_dioxane_on_special_position
15. GIHW0J_mol_on_special_position
16. GUCSEC_solvent_on_special_position
17. HIHF EK_water_on_special_position
18. JINGIW01_chloride_on_special_position
19. JOHFUJ_mol_on_special_position
20. JUBFER_molecule_on_special_position
21. JUKWAN_molecule_on_special_position
22. KADKIJ_on_special_position
23. KANXIG_h_molecule_on_special_position
24. KHDBEZ_h_K_on_special_position
25. LCYSTI14_molecule_on_special_position
26. LIKVEI_acn_on_special_position
27. LOYCUZ_solvent_on_special_position
28. MAAZNB_h-molecule_on_special_position
29. MAVGAS_counterion_on_special_position
30. NEQVEL_mol_on_special_position
31. NOJKON_cofomer_on_special_position
32. OGEREY_sulphate_on_special_position
33. OGUTOB_cofomer_on_special_position
34. OJUQAN_cofomer_on_special_position
35. ORACUH_solvent_on_special_position
36. QIGCEO05_molecule_on_special_position
37. QUOLSO_on_special_position
38. RIGTOS01_molecule_on_special_position
39. SALHIW_chloride_on_special_position
40. SIMJUU_mol_on_special_position
41. SIXZOO02_molecule_on_special_position
42. TIWXOM01_molecule_on_special_position
43. VIKLUX_fumaric_a_on_special_position

44. WAXZEC_cofomer_on_special_position
45. WECGUI_water_on_special_position
46. XOGDEE_solvent_on_special_position
47. XOVBIU_molecule_on_special_position
48. YAWROF_molecule_on_special_position
49. YOWTAH01_molecule_on_special_position
50. YOYSUC_molecule_on_special_position
51. ZAZLER_cofomer_on_special_position
52. ZZZDRQ01_molecule_on_special_position
53. ZZZVZA05_molecule_on_special_position

Table 2: List of 24 disordered structures contained in the Drug Subset that were optimized using two conformers. These were thus present twice, denoted REFCODE_1 and REFCODE_2 in the following list of 732 optimized structures.

1. BEQGIN
2. BULCUG03
3. DIKWEA
4. DULKAX
5. FMDURD02
6. HAFNOR
7. IHOZOW
8. ILUSAL
9. JOWCAB02
10. KABZES
11. LCYSTN28
12. LERJID
13. MIFZIK
14. MUCDUK
15. NIQVIT
16. NUYSEG
17. PCLPMS
18. SEBPEU01
19. SIWCEG
20. SOYMID
21. TADZAZ03
22. VIGDAR
23. WETPOB
24. YIGPIO02

Table 3: List of structures where optimization failed or was not performed due to missing atoms, disorder, and/or combined with structural complexity.

1. ACXMAC
2. ADENTP³
3. AMASIS
4. BAGVUA
5. CEFMEN_h
6. DAHREJ
7. FUFJUL⁴
8. IDIVOG⁵
9. JINGIW01
10. LUQREW
11. LUXYOU⁶
12. NAQLAU⁷
13. NUYSEG
14. NACGLY03_h
15. OGIZEJ⁸
16. RUWHEX
17. SUCHOL12
18. SUDXAQ
19. TEZZUV_h
20. TMTUBC_h
21. TUCMEJ⁹
22. TUSHIZ¹⁰
23. YICMUS¹¹
24. ZEFNOP

Table 4: List of 708 optimized structures. 732 structures are listed, because the 24 disordered structures appear twice, one for each optimized conformer.

1. ABABIQ_r
2. ACEROR
3. ACHOLC
4. ACPRET03
5. ACSALA14
6. ACTDGU10_h
7. ADENOS10
8. ADPOSM01_h
9. ADRTAR
10. AHILOU
11. AIPBAR_h
12. AKENOU
13. AKOFUD
14. AMAVEP
15. AMAVIT
16. AMAVOZ
17. AMBZCL
18. AMCAPR11
19. AMEQAK
20. AMEVET
21. AMMCHC11
22. AMOXCT10_h
23. AMPCIH01
24. AMYTAL10_h
25. ANIXAX
26. ANSTER10
27. ANTPYR
28. APUYOA02
29. ARFCYT10
30. ARFUAD
31. ARISOK_h
32. ASPARM06
33. ASPART10
34. ATDZSA03
35. AXAKOA
36. BAFVOT
37. BAHDET
38. BASCUT_h
39. BEDLUR
40. BEDMIG04
41. BENWUO
42. BEQGIN_1
43. BEQGIN_2
44. BEWKUJ15
45. BEZJIA

46. BIDFUQ
47. BIHGUV
48. BIHYEW10
49. BIKHIN01
50. BILJEN
51. BIOTIN12
52. BIRKES10
53. BISMEV11
54. BIXFOD_h
55. BIXGIY03_h
56. BIYSEH07
57. BOCYAV
58. BOLBOV
59. BOMDUC
60. BOVDUM
61. BPYZDO21_h
62. BUBYII
63. BUFPAV
64. BUGJOC
65. BUHCIQ_h
66. BULCUG03_1
67. BULCUG03_2
68. BZPENK01
69. BZTRMS_h
70. CABCU06
71. CAGWOW
72. CAGWUC_h
73. CAGXUD
74. CAKYOE
75. CALDAU_h
76. CANDOM10_h
77. CANDUR
78. CANFED
79. CARPEQ_h
80. CATBOP
81. CAVVUQ02
82. CEKBEZ
83. CELCIF
84. CEPGLY_h
85. CEPHHM10_h
86. CETHNA_h
87. CEYHUJ01_h
88. CFHPRG
89. CHEBAR01
90. CIDDEZ
91. CIHHIK01
92. CIMETD02
93. CINSEZ

94. CIPWUT_h
95. CIRTIG01_h
96. CISKUL
97. CITREC01
98. CITYIN10
99. CIVFOC10
100. CIYRIL_h
101. CLAMBU
102. CLDZPA01
103. CLEMAS02
104. CLMPCL02
105. CLONDC11
106. CLPNXA
107. COCHCL01
108. COQWOU02
109. CORYIR10_h
110. COTYEP
111. COTYOA02
112. COTYOA
113. COVPIN03
114. COXYOE
115. COYSAK
116. CPHNZP
117. CPPBGC
118. CPTAZP_h
119. CPTZAP_h
120. CREATH01
121. CUBHUE
122. CUYCEF01
123. CUYDOQ_w_h
124. CXMCYT
125. CYPROT10
126. DABJAR
127. DACMOR01
128. DAPSUO05
129. DAQCII
130. DARGOT03
131. DAUNMY_h
132. DAXYUX01_h
133. DBEZPO01_h
134. DEFZEU
135. DEMXUO
136. DEQYIH
137. DESNAR
138. DEWHOC
139. DEXMET11
140. DEXPOX_h
141. DFIMZP01

142. DIAZOX
143. DIGOXN10
144. DIJWOJ
145. DIKWEA_1
146. DIKWEA_2
147. DINXIJ
148. DIRCIS
149. DIVHOF02_h
150. DIWQIJ
151. DIYYOB
152. DIZPAM11
153. DIZWAK_h
154. DLPROM02
155. DLSERN31
156. DMANTL07
157. DMTYRS_h
158. DOLJIX_h
159. DOQBUI
160. DPHPZL
161. DPYALC
162. DUDGAL
163. DUJFAQ
164. DUJPOO
165. DUKTOS_h
166. DULKAX_1
167. DULKAX_2
168. DUNGAV
169. DUNXAL_h
170. DUTRUF10
171. DUVZOJ02_h
172. DXMTAS
173. EDALEC_m
174. EFAXAM
175. EFEHOO
176. EFEMUX01
177. EGINEP
178. EHOXEE_h
179. EJEQAL
180. ELIZAC
181. ELOVAE01
182. ELOVOS
183. EMIQEY
184. ENEBOQ
185. EPHPMO
186. EQONOP
187. ESOURCE11
188. ESTRIO
189. ESTRON14

190. ETBBAR02
191. ETCYPY01
192. ETEXIK02
193. ETHACN_h
194. ETHDPH05
195. ETHUSS03
196. ETPRGN_h
197. EVAFUD
198. EXAQEA
199. EXAQIE
200. EXAQOK
201. FABVAF01
202. FAHFVAV01
203. FAHMAC
204. FAJKUW
205. FAJMAE
206. FANNEN_h
207. FARRUM02
208. FAXYAG
209. FDOURD01
210. FEBHUP02
211. FEFNEK
212. FEMBII
213. FETGAN
214. FHMANE
215. FIDYIA01
216. FIFDON_h
217. FIKFIP
218. FILGEM03_h
219. FIMNIY
220. FIMTAW
221. FINWEE_h
222. FISLIE
223. FITPIH02
224. FIXGAU01
225. FIXNUV
226. FIZWAM_h
227. FMDURD02_1
228. FMDURD02_2
229. FOGVIG02
230. FOHLIY01
231. FOLCAH03
232. FOPYUE_h
233. FOQREK
234. FORGAU01
235. FOSPEA
236. FOYMAH
237. FPRTOD10

238. FUFTIK
239. FUGSEF
240. FUHGOE
241. FUMKEF
242. FUPDOJ
243. FUPFAX_h
244. FUQMOU01
245. FUWNIU
246. FUXMOA01
247. GABFOE_h
248. GAHHOM01
249. GAKWIY_h
250. GALDEC01
251. GAMFAC
252. GAQTAU01
253. GARFEK
254. GEGJAD
255. GEHXEX
256. GERXAE
257. GICHEG
258. GICVUJ01_h
259. GIDLUB
260. GILHAL
261. GIMYEG
262. GIXXOB
263. GLCROL01
264. GLUCIT
265. GLUTAM02
266. GLUTAS05
267. GLYCIN99
268. GODTIC
269. GOHSUS
270. GOLWIN
271. GOPDUK01
272. GORKUV01
273. GRISFL02
274. GUBLAS
275. GUNKEH
276. GUYGOX03_h
277. HAFNOR_1
278. HAFNOR_2
279. HAKJIM
280. HANEDP10
281. HAXHET_h
282. HCCYHG_h
283. HEPHCL10
284. HERGOM_h
285. HEYGEX

286. HIJFAH
287. HISHEX
288. HISTPA15
289. HMXBZP
290. HOQHUQ
291. HOTTER
292. HUCHEU02
293. HUHZEP
294. HXVITD
295. HYDLAZ01
296. IBUNOJ
297. ICRFRB10
298. IDOXUR01
299. IFOSAY
300. IGAZUM_h
301. IHOZOW_1
302. IHOZOW_2
303. IHUHAW_h
304. IKACUT
305. ILUSAL_1
306. ILUSAL_2
307. IMEHOY
308. INAHZC_h
309. INOMUV
310. IPILUQ
311. IPMEPL
312. IPRAMI_h
313. IQILAW
314. IQISAE
315. IQUMAI
316. IVEGIA_h
317. IVUQIZ05
318. IWOCON
319. JAKGEH
320. JAMRIY
321. JAVCIS10
322. JAZCET
323. JEBNUZ_h
324. JIWPEL01
325. JIWQAI03
326. JODZUX
327. JOHKUM
328. JOPSUE
329. JORJUX
330. JOSWAP01_h
331. JOWCAB02_1
332. JOWCAB02_2
333. JUPDAB

334. JUYLEW
335. JUYZOS
336. KABGUR
337. KABZES_1
338. KABZES_2
339. KAGCIE
340. KASMOH
341. KAXSEJ
342. KCANRO
343. KEBGAB01
344. KEBVES_h
345. KEHBAA
346. KEMZIL01
347. KEYSOW
348. KIDRAR01
349. KIDWAW
350. KIJMEX
351. KIKWUW
352. KIMDER
353. KIPLEC
354. KIWQUC
355. KNMYSL
356. KOFNIC
357. KORSAL02_h
358. KORYUL
359. KUGWIS_h
360. KUQDAC01_h
361. KURJAK
362. KUTJIU
363. LADTIT
364. LAKZII
365. LALNIN03
366. LANLET
367. LAQBAI
368. LAQSON01
369. LARGPH06
370. LATPIH
371. LATQAA
372. LATSOQ01_h
373. LAYCET01
374. LCYSTN28_1
375. LCYSTN28_2
376. LDOPAS03
377. LEBLUB
378. LERJID_1
379. LERJID_2
380. LESVEM
381. LEXQIQ

382. LEZREO_h
383. LINTUX_h
384. LOPZEX
385. LORCAW_h
386. LOSLEL
387. LOWKOX_h
388. LTXPSE_h
389. LUDDUJ
390. MACKUX
391. MACXPR10
392. MAFLAI_h
393. MCPRPL01
394. MELATN01
395. MELFIT09
396. MEQCOB
397. METHCL01
398. MEXSEP
399. MEYGII
400. MHOCUM01
401. MIFPOI
402. MIFZIK_1
403. MIFZIK_2
404. MIGQID
405. MIHZUA
406. MIMOSN_h
407. MIRACD_h
408. MNIMET02
409. MOKXIU
410. MOKXOA
411. MOPHAC
412. MORPHM01
413. MOSXUO
414. MOVLOZ
415. MOVYAY01_h
416. MOXAZO_h
417. MPIXPS_h
418. MPZPAM
419. MTEPAM
420. MTHPRG
421. MUBPAB
422. MUCDUK_1
423. MUCDUK_2
424. MUGKAA01
425. MUYVAE_h
426. MVERIQ01
427. NADREN
428. NAIPOE
429. NAKGOY

430. NALCYS10
431. NAMDPC01_m
432. NANNUN
433. NAPHZL10_h
434. NARBOY
435. NAZLAC02
436. NDNHCL01
437. NEPHCL
438. NEQGUM
439. NEQHEY
440. NESKED
441. NETIND01
442. NEVMOR
443. NHPBZO_h
444. NICOAM05
445. NIHMOH01
446. NIKROP
447. NILXOV_m
448. NIQVIT_1
449. NIQVIT_2
450. NIRRAH_h
451. NIVBIO_h
452. NOBDUF
453. NOKGAX
454. NORGES01
455. NORMET_h
456. NUJDAX_w
457. NUPQEW
458. NUYHUK
459. NUYSEG_1
460. NUYSEG_2
461. OBEQAN01
462. OCAHAC
463. OFAZUQ
464. OFEJOA
465. OFILEW
466. OGIXIN
467. OMAKOE01
468. OPOGAD
469. OPOQER
470. ORNHCL12
471. OROZAX
472. OXYTET01_h
473. PABHIJ01
474. PAJNUJ
475. PANCUR
476. PBUMSL_h
477. PBUMSL

478. PCLPMS_1
479. PCLPMS_2
480. PEBGUC02
481. PEDWOM
482. PEKFAN
483. PERPAZ
484. PETHID_h
485. PEXKOS01
486. PEYGOQ
487. PGDEDO
488. PHEPHR
489. PICVUU01
490. PIGSOO
491. PILOCP01
492. PIMKOL
493. PIMOZD
494. PINDON
495. PIPAMP
496. PIPCIL_h
497. PIRENA10_w
498. PIWLEM02
499. PMEPEN01
500. POBKOG
501. POFORM_m
502. POGKOM
503. PRAZAM_h
504. PRGDOL02
505. PROCPH
506. PROGLE20_h
507. PROGST12
508. PUFGUU
509. PUQGUF
510. PUVMAU
511. PUVRIH
512. PYRXCL01
513. PYRZIN23
514. QABCII
515. QABTOD01
516. QATNEF01
517. QECHOV03
518. QEMLIE
519. QETZUL
520. QICMEU_h
521. QIFKEX01
522. QIJPAC
523. QIJZOY
524. QIQKEI
525. QIQLIL

526. QOMKUA
527. QQQAEJ02
528. QQQAUG04
529. QQQBSS01_h
530. QUBQIO01
531. QURWOQ02
532. QUZWOY
533. RALMIA
534. RALMOG01
535. RASTAG_m
536. RAVPUB
537. RBFLAV10_h
538. REBCIL
539. REDNIX
540. REZBII
541. RIGDES
542. RILQOU
543. RINTIT
544. RIQFOO
545. RIWKOX
546. ROKNUB
547. ROQLOA
548. RSERPNO1
549. RUKBOP
550. RUKQOF
551. RUTGOE
552. RUTGUK
553. RUWBAM
554. RUYZAO
555. SAFNIW
556. SAHSOJ_h
557. SAJFIT
558. SAJNOG_h
559. SAQYIR
560. SATHOL
561. SAWFIF_h
562. SAXFED
563. SAXQIT
564. SAZLAG
565. SAZNOW_m
566. SEBPEU01_1
567. SEBPEU01_2
568. SEBPEU01
569. SENJIF
570. SEQKAA02
571. SEQNIL
572. SESKUY01
573. SETSOA

574. SFDMOX05_h
575. SIFGOD
576. SIJWUF
577. SIKLIH07
578. SIWCEG_1
579. SIWCEG_2
580. SLFNMB02
581. SLFNMC20_h
582. SLFNMG01
583. SLFSXZ12
584. SOBBUG_h
585. SONKEM
586. SOPSEV
587. SOYMID_1
588. SOYMID_2
589. SUJXUR
590. SUTHAZ37
591. SUTWIO
592. SUVGUL
593. SUWNUT
594. TADLIU02
595. TADZAZ03_1
596. TADZAZ03_2
597. TAJFOZ_h
598. TANDET
599. TAYFUV_h
600. TAZZUP
601. TEPHAM_h
602. TESTON10
603. TETCYH12
604. TEZZEE
605. TFPROM10
606. THCLCS_h
607. THGUAN10
608. TIJMII
609. TIPWAQ_h
610. TITHYN10_h
611. TIYQAU01
612. TMPHCL01
613. TOKSES_h
614. TOPXED_m
615. TOXXIN_h
616. TOYJOG
617. TOYSUX
618. TRHTRT_h
619. TUBOBM_h
620. TULRIB_m
621. TUQGET

622. TUTDOD
623. TUZCOI
624. UCUVET
625. UGIYEP
626. UHUQUK
627. UKAKIA
628. UKIFIE_h
629. UMOMAL
630. UPUKOH
631. UQAPEJ
632. URAVUG
633. UREA0H12
634. USOCOV
635. UXIXUV
636. VAMBOA01
637. VAWPIT
638. VAWQEQ
639. VAYJOW
640. VAYXOI
641. VEMRAI
642. VEVQAQ
643. VICXOV
644. VIDJEX01
645. VIFQIL
646. VIGDAR_1
647. VIGDAR_2
648. VOGCAV_h
649. VOXDUH
650. VUCDUS
651. VUCJEI
652. VUJRUO_h
653. VUXPUZ
654. WACHIR_h
655. WADPIC
656. WADTIG
657. WAHJAR
658. WAJYUC
659. WALPIJ
660. WASGAA_h
661. WAWVOH
662. WEHYEO
663. WELGAV
664. WEMGEC
665. WEQHEF01_h
666. WERVEU02
667. WETLOY
668. WETPOB_1
669. WETPOB_2

670. WEXQAS01
671. WEZCOT
672. WIBZEM
673. WINJAE_m
674. WINWUL02
675. WIPYOL
676. WOQDAH
677. WOYPAB
678. WURPUU
679. WUTHEA01
680. XANTOX_h
681. XAPTAK01
682. XAQNOU
683. XATJAF
684. XAVTOF02
685. XAYGEJ01
686. XAZRUN
687. XEJKUS
688. XEZTON01
689. XICRUY
690. XIJKIK01
691. XIRHOX
692. XOCXUI01
693. XOGMOX
694. XOJNOB
695. XOSGUH
696. XOZRUZ
697. XULRUT
698. YACGUE_h
699. YACHAL_h
700. YACTEC01
701. YALWIS
702. YALYUH
703. YAPZEU
704. YARXEW
705. YIGPIO02_1
706. YIGPIO02_2
707. YIGRAI
708. YIKNUD
709. YIYRAZ
710. YODXAR
711. YOKQAQ
712. YUKVAD
713. YURSUB_h
714. ZAPLEJ
715. ZAPXOF
716. ZEMJOQ
717. ZENREP

718.	ZIDLED
719.	ZIKJOS
720.	ZIMBOO
721.	ZIVKEU
722.	ZIZRUX01
723.	ZOGSOD
724.	ZOXSEK
725.	ZUHFIR
726.	ZUQMIJ_h
727.	ZZZECG01
728.	ZZZPNG01
729.	ZZZPUS19
730.	ZZZRCG01_h
731.	ZZZSHG01
732.	ZZZTSE03

refcode	rmsd		Tanimoto_rmsd
ABABIQ_ro		0.10962	0.99925
ACERORo		0.05292	0.99956
ACHOLCo		0.14007	0.99662
ACPRET03o		0.06479	0.99975
ACSALA14o		0.03692	0.99975
ACTDGU10_ho		0.19352	0.99882
ADENOS10o		0.04381	0.9998
ADPOSM01_ho		1.29284	0.86729
ADRTARo		0.09257	0.99897
AHILOUo		0.16949	0.99922
AIPBAR_ho		0.08951	0.99859
AKENOUo		0.18442	0.9992
AKOFUDo		0.08304	0.99952
AMAVEPo		0.07934	0.99939
AMAVITo		0.06364	0.99952
AMAVOZo		0.07694	0.99961
AMBZCLo		0.06669	0.99966
AMCAPR11o		0.12386	0.99818
AMEQAKo		0.04025	0.9999
AMEVETo		0.08915	0.99963
AMMCHC11o		0.11217	0.99804
AMOXCT10_ho		0.11275	0.99918
AMPCIH01o		0.0806	0.99957
AMYTAL10_ho		0.04831	0.99968
ANIXAXo		0.05647	0.99966
ANSTER10o		0.04046	0.99986
ANTPYRo		0.05262	0.99958
APUYOAO2o		0.09924	0.99951
ARFCYT10o		0.05835	0.99958
ARFUADo		0.07592	0.99936
ASPARM06o		0.05009	0.99932
ASPART10o	Initially two hydrogen atoms missing, corrected		n/a
ATDZSA03o		0.07522	0.99932
AXAKOAO		0.1522	0.99817
BAFVOTo		0.03027	0.99984
BAHDETo		0.141	0.99835
BASCUT_ho		0.12558	0.99771
BEDLURo		0.07748	0.99973
BEDMIG04o		0.22336	0.99477
BENWUOo		0.04949	0.99977
BEQGIN_1o		0.12694	0.999
BEQGIN_2o		0.35614	0.99218
BEWKUJ15o		0.06383	0.99954
BEZJIAo		0.06415	0.99969
BIDFUQo		0.07285	0.99984
BIHGUVo		0.16461	0.99736
BIHYEW10o		0.06408	0.99946
BIKHIN01o		0.02317	0.99985
BILJENo		0.1137	0.99888

BIOTIN12o	0.06949	0.9996
BIRKES10o	0.16381	0.9985
BISMEV11o	0.04256	0.99959
BIXFOD_ho	0.13603	0.99784
BIXGIY03_ho	0.04319	0.99981
BIYSEH07o	0.08263	0.9995
BOCYAVo	0.08305	0.99951
BOLBOVo	0.09515	0.99982
BOMDUCo	0.077	0.99955
BOVDUMo	0.29245	0.99613
BPYZDO21_ho	0.12853	0.9985
BUBYIlo	0.05154	0.99965
BUFPAVo	0.10324	0.99892
BUGJOCo	0.04629	0.99985
BUHCIQ_ho	0.08469	0.99936
BULCUG03_1o	0.02467	0.99977
BULCUG03_2o	0.02469	0.99977
BULCUG03_3o	0.02482	0.99977
BULCUG03_4o	0.02461	0.99977
BULCUG03_5o	0.02368	0.99979
BULCUG03_6o	0.02401	0.99979
BULCUG03_7o	0.02402	0.99979
BULCUG03_8o	0.02402	0.99979
BZTRMS_ho	0.10244	0.99912
CABCUD06o	0.08292	0.99955
CAGWOWo	0.07614	0.99937
CAGWUC_ho	1.78426	0.73287
CAGXUDo	0.0887	0.99931
CAKYOEo	0.04337	0.99982
CALDAU_ho	0.11941	0.99946
CANDOM10_ho	0.11674	0.99914
CANDURo	0.07844	0.99919
CANFEDo	0.07756	0.99938
CARPEQ_ho	0.43989	0.99003
CATBOPo	0.07741	0.9994
CAVVUQ02o	0.57497	0.96508
CEKBEZo	0.0827	0.99958
CELCIFo	0.10233	0.99961
CEPGLY_ho	0.25241	0.99651
CEPHHM10_ho	0.11777	0.9993

Considerable changes in optimized structure. Input structure without H. Ion shifted considerably. OK

CETHNA_ho		n/a
CFHPRGo	0.13751	0.99874
CHEBAR01o	0.06957	0.99936
CIDDEZo	0.07596	0.99908
CIHHIK01o	0.13656	0.99844
CIMETD02o	0.12948	0.9985
CINSEZo	0.103	0.99948
CIPWUT_ho	0.10846	0.99881
CIRTIG01_ho	0.27882	0.99536

CISKULo		0.22902	0.99825
CITYIN10o		0.10905	0.99888
CIVFOC10o		0.05992	0.99944
CIYRIL_ho		0.04875	0.9998
CLAMBUo		0.15218	0.99837
CLDZPA01o		0.10356	0.99883
CLEMAS02o		0.14308	0.99845
CLMPCL02o		0.08617	0.9993
CLONDC11o		0.12006	0.99792
CLPNXAo		0.03572	0.99994
COCHCL01o		0.06109	0.99969
COQWOU02o		0.09949	0.99934
CORYIR10_ho		0.09474	0.99895
COTYEPo		0.0406	0.99978
COTYOA02o		0.11046	0.99806
COTYOAo		0.13901	0.99807
COVPIN03o		0.09276	0.99938
COXYOEo		0.12382	0.99864
COYSAKo		0.05711	0.99969
CPHNZPo		0.09412	0.99913
CPPBGCo		0.22256	0.99566
CPTAZP_ho		1.38006	0.92718
CPTZAP_ho		0.08739	0.99924
CREATH01o		0.08448	0.99819
CUBHUEo		0.06637	0.99934
CUYCEF01o		0.08657	0.99948
CUYDOQ_w_ho		0.04818	0.9997
CXMCYTo		0.0866	0.99912
CYPROT10o		0.04715	0.99985
DABJARo		0.07619	0.99959
DACMOR01o		0.26037	0.99386
DAPSUO05o		0.05112	0.99973
DAQCIIo		0.069	0.99971
DARGOT03o		0.08415	0.99971
DAUNMY_ho		0.14928	0.99895
DAXYUX01_ho		0.16565	0.99835
DBEZPO01_ho		0.06578	0.99968
DEFZEUo		0.08982	0.99946
DEMXUOo		0.08621	0.99882
DEQYIHo		0.04534	0.9998
DESNARo	0.131, OK		
DEWHOCo		0.04765	0.99976
DEXMET11o		0.07047	0.99964
DEXPOX_ho		0.15367	0.99794
DFIMZP01o		0.07801	0.99939
DIAZOXo		0.03537	0.99981
DIGOXN10o		0.12445	0.99977
DIJWOJo		0.12776	0.99942
DINXIJo		0.21488	0.99885
DIRCISo		0.12426	0.99935

DIVHOF02_ho	0.61576	0.97992
DIWQIJJo	0.11525	0.99955
DIYYOBo	0.0617	0.99953
DIZPAM11o	0.0451	0.99979
DIZWAK_ho	0.06928	0.99973
DLPROM02o	0.04439	0.9994
DLSERN31o	0.08911	0.99712
DMANTL07o	0.0237	0.99991
DMTYRS_ho	0.04613	0.99969
DOLJIX_ho	0.34664	0.98908
DOQBUIo	0.0951	0.9995
DPHPZLo	0.1102	0.99879
DPYALCo	0.17146	0.99758
DUDGALo	0.05584	0.9997
DUJFAQo	0.15444	0.99903
DUKTOS_ho	1.61333	0.81462
DULKAX_1o	0.11129	0.99945
DULKAX_2o	0.26958	0.99677
DUNGAVo	0.08791	0.99926
DUNXAL_ho	0.15772	0.99927
DUTRUF10o	0.11135	0.99943
DUVZOJ02_ho	0.11271	0.99845
DXMTASo	0.09835	0.99931
EDALEC_mo	P-XRD structure with considerable differences to optimized one, OK n/a	
EFAXAMo	0.13809	0.99973
EFEHOOo	0.07661	0.99979
EFEMUX01o	0.04681	0.99973
EGINEPo	0.09409	0.99945
EHOXEE_ho	0.13647	0.99865
EJEQALo	0.10494	0.99933
ELIZACo	0.10226	0.99918
ELOVAE01o	0.02879	0.99989
ELOVOSo	0.05711	0.99986
EMIQEYo	0.2706	0.99666
ENEBOQo	0.15034	0.9975
EPHPMOo	0.06313	0.99935
EQONOPo	0.23767	0.99532
ESOURE11o	0.02618	0.99994
ESTRIOo	0.05159	0.99978
ESTRON14o	0.05059	0.99977
ETBBAR02o	0.07399	0.99919
ETCYPY01o	0.04293	0.99968
ETEXIK02o	0.08854	0.99925
ETHACN_ho	0.73812	0.96026
ETHDPH05o	0.12333	0.99665
ETHUSS03o	0.06435	0.99965
ETPRGN_ho	0.56779	0.97142
EVAFUDo	0.04249	0.99983
EXAQEAo	0.10733	0.99908
EXAQIEo	0.05257	0.99972

EXAQOKo		0.04498	0.9998
FABVAF01o		0.15763	0.99829
FAHFAV01o		0.10849	0.9992
FAHMACo		0.10082	0.99926
FAJKUWo		0.16041	0.99864
FAJMAEo		0.13122	0.99893
FANNEN_ho	0.0853, OK		n/a
FARRUM02o		0.05746	0.99985
FAXYAGo		0.19805	0.99821
FDOURD01o		0.06256	0.99956
FEBHUP02o		0.047	0.99988
FEFNEKo		0.0782	0.99954
FEMBIlo		0.11881	0.9994
FETGANo		0.07326	0.99981
FHMANEo		0.05817	0.99969
FIFDON_ho		0.06325	0.99976
FIKFIPo		0.06422	0.99955
FILGEM03_ho		0.05022	0.99972
FIMNIYo		0.16089	0.99756
FIMTAWo		0.10236	0.99905
FINWEE_ho		0.18695	0.99874
FISLIEo		0.21804	0.99415
FITPIH02o		0.02886	0.99984
FIXGAU01o		0.07751	0.99942
FIXNUVo		0.30538	0.99066
FIZWAM_ho		2.1554	0.81824
FMDURD02_1o		0.09604	0.99903
FMDURD02_2o		0.46354	0.97838
FOGVIG02o		0.0862	0.99924
FOHLIY01o		0.05938	0.9995
FOLCAH03o		0.16186	0.99918
FOPYUE_ho		0.14679	0.99949
FOQREKo		0.03568	0.99985
FORGAU01o		0.08504	0.99952
FOSPEAo		0.09092	0.99767
FOYMAHo		0.0536	0.99928
FPRTOD10o		0.07823	0.99955
FUFTIKo		0.08343	0.99892
FUGSEFo		0.09305	0.99927
FUHGOEo		0.07015	0.99957
FUMKEFo		0.07401	0.99953
FUPDOJo		0.08922	0.99958
FUQMOU01o		0.24981	0.99438
FUWNIUo		0.10236	0.9988
FUXMOA01o		0.04601	0.99983
GABFOE_ho		0.30211	0.9963
GAKWIY_ho		0.12795	0.99926
GALDEC01o		0.02062	0.99989
GAMFACo		0.09087	0.99907
GAQTAU01o		0.02325	0.99993

GARFEKo	0.13618	0.99857
GEHXEXo	0.03755	0.99986
GERXAEo	0.19917	0.99801
GICHEGo	0.34667	0.99164
GICVUJ01_ho	0.07365	0.99977
GIDLUBo	0.02568	0.99987
GILHALo	0.0747	0.99979
GIMYEGo	0.10895	0.99963
GIXXOBo	0.06733	0.99986
GLCROL01o	0.01897	0.99988
GLUCITo	0.24703	0.99105
GLUTAM02o	0.15289	0.99577
GLUTAS05o	0.18215	0.99777
GLYCIN99o	0.14877	0.98915
GODTICo	0.0445	0.99982
GOHSUSo	0.0917	0.99934
GOLWINo	0.0422	0.99989
GOPDUK01o	0.16992	0.99851
GORKUV01o	0.58293	0.9524
GRISFL02o	0.07489	0.9995
GUBLASo	0.15622	0.99878
GUNKEHo	0.06835	0.99976
GUYGOX03_ho	0.2156	0.99652
HAFNOR_1o	0.30616	0.99625
HAFNOR_2o	0.36496	0.99469
HAKJIM_ho	H was initially missing. Corrected.	n/a
HANEDP10o	0.08547	0.99937
HAXHET_ho	0.11878	0.99948
HCCYHG_ho	0.11827	0.99735
HEPHCL10o	0.03082	0.99988
HERGOM_ho	0.12258	0.99945
HEYGEXo	0.03909	0.99978
HIJFAHo	0.1539	0.99919
HISHEXo	0.06325	0.99958
HISTPA15o	0.04951	0.99962
HMXBZPo	0.04678	0.99979
HOQHUQo	0.01123	0.99994
HOTTERo	0.08056	0.99939
HUCHEU02o	0.12968	0.99947
HUHZEPo	0.06862	0.9995
HXVITDo	0.04126	0.99994
HYDLAZ01o	0.0484	0.99955
IBUNOJo	0.05303	0.9997
ICRFRB10o	0.04168	0.99987
IDOXUR01o	0.05415	0.99967
IFOSAYo	0.06216	0.99957
IHOZOW_1o	0.21223	0.99743
IHOZOW_2o	0.32726	0.99394
IHUHAW_ho	0.26596	0.99691
IKACUTo	0.02813	0.99984

ILUSAL_1o		0.07959	0.99962
IMEHOYo		0.08374	0.99922
INAHZC_ho		0.07816	0.99878
INOMUVo		0.06678	0.99967
IPILUQo		0.07306	0.99963
IPMEPLo		0.01962	0.99993
IPRAMI_ho		0.13013	0.99856
IQILAWo		0.07373	0.99981
IQISAEo		0.18946	0.99777
IQUMAIo		0.11398	0.99908
IVUQIZ05o		0.06183	0.99964
IWOCONo		0.06021	0.99976
JAKGEHo		0.06924	0.9996
JAMRIYo		0.06701	0.99886
JAZCETo		0.35546	0.99744
JEBNUZ_ho		0.23532	0.9968
JIWPEL01o		0.11087	0.99912
JIWQAI03o		0.07602	0.99961
JODZUXo		0.39612	0.99392
JOHKUMo		0.13805	0.99907
JOPSUEo		0.05955	0.99956
JORJUXo		0.08261	0.99942
JOSWAP01_ho		0.11319	0.99913
JOWCAB02_1o		0.10776	0.9996
JOWCAB02_2o		0.25153	0.99784
JUPDABo		0.06133	0.99934
JUYLEWo		0.03238	0.99987
JUYZOSo		0.07788	0.99952
KABGURo		0.0979	0.99901
KABZES_1o		0.23063	0.9959
KABZES_2o		0.23489	0.99574
KAGCIEo		0.14225	0.99896
KASMOHo		0.0831	0.9992
KAXSEJo		0.03113	0.9999
KEBGAB01o		0.08246	0.99864
KEBVES_ho		0.63653	0.98137
KEHBAAo		0.15385	0.99903
KEMZIL01o		0.00245	0.99998
KEYSOWo		0.20762	0.99651
KIDRAR01o		0.07301	0.99964
KIDWAWo		0.14639	0.99876
KIJMEXo		0.08174	0.99949
KIKWUWo		0.13292	0.99887
KIMDERo		0.11794	0.99939
KIPLECo		0.11329	0.99933
KIWQUCo	0.162, OK		n/a
KNMYSLo		0.13311	0.99912
KOFNICo		0.09246	0.99903
KORSAL02_ho		0.06568	0.9981
KORYULo		0.04068	0.99984

KUGWIS_ho	0.08963	0.99925
KUQDAC01_ho	0.1885	0.99785
KURJAKo	0.10878	0.9978
KUTJIUo	0.08488	0.99933
LADTITo	0.03506	0.99984
LAKZIIo	0.18684	0.99841
LALNIN03o	0.10104	0.99573
LANLETo	0.06283	0.99966
LAQBAlO	0.04132	0.99984
LAQSON01o	0.13359	0.99921
LARGPH06o	0.12111	0.99816
LATPIHo	0.11703	0.99898
LATQA Ao	0.10556	0.99925
LAYCET01o	0.05042	0.99974
LCYSTN28_1o	0.11336	0.9957
LCYSTN28_2o	0.03481	0.99959
LDOPAS03o	0.04774	0.99975
LESVEMo	0.07517	0.99978
LEXQIQo	0.0951	0.99949
LINTUX_ho	0.03153	0.99988
LOPZEXo	0.25765	0.9973
LORCAW_ho	0.06848	0.99958
LTXPSE_ho	0.10068	0.99883
LUDDUJo	0.17077	0.99745
MACKUXo	0.09777	0.99951
MACXPR10o	0.06703	0.99969
MCPRPL01o	0.06725	0.99936
MELATN01o	0.06661	0.99966
MELFIT09o	0.19868	0.99899
MEQCOBo	0.09652	0.99973
METHCL01o	0.15682	0.99557
MEXSEPo	0.03772	0.99986
MEYGIIo	0.11699	0.99808
MHOCUM01o	0.09383	0.99938
MIFPOIo	0.02687	0.99994
MIFZIK_1o	0.16046	0.99618
MIFZIK_2o	0.17576	0.99543
MIGQIDo	0.06523	0.99955
MIHZUAo	0.09869	0.99935
MIMOSN_ho	0.06378	0.99953
MIRACD_ho	0.48916	0.98407
MNIMET02o	0.07684	0.99879
MOKXIUo	0.1871	0.99858
MOKXOAo	0.08621	0.99959
MOPHACo	0.06137	0.99935
MORPHM01o	0.0489	0.99968
MOSXUOo	0.02285	0.99991
MOVLOZo	0.27855	0.97669
MOVYAY01_ho	0.182	0.99844
MOXAZO_ho	0.06206	0.99896

MPIXPS_ho	0.503, molecule was shifted, OK	n/a	
MPZPAMo		0.04426	0.99983
MTEPAMo		0.07018	0.99975
MTHPRGo		0.06835	0.99967
MUBPABo		0.07506	0.99956
MUCDUK_1o		0.11722	0.99888
MUCDUK_2o		0.89743	0.93625
MUGKAA01o		0.04067	0.99975
MUYVAE_ho		0.11963	0.99838
MVERIQ01o		0.06224	0.99971
NADRENo		0.02977	0.99987
NAJPOEo		0.1776	0.99881
NAKGOYo		0.04871	0.99956
NALCYS10o		0.11783	0.99719
NAMDPC01_mo	0.0249, OK	n/a	
NANNUNo		0.08421	0.99934
NARBOYo		0.16148	0.99692
NAZLAC02o		0.1739	0.99891
NDNHCL01o		0.08564	0.9994
NEPHCLo		0.03218	0.99979
NEQGUMo		0.15798	0.99872
NEQHEYo		0.1168	0.99932
NESKEDo		0.07268	0.99958
NETIND01o		0.04863	0.99981
NEVMORo		0.0407	0.99973
NHPBZO_ho		0.11397	0.99874
NICOAM05o		0.0205	0.99989
NIHMOH01o		0.1863	0.99801
NIKROPo		0.12923	0.99824
NILXOV_mo	Molecules shifted into unit cell, OK	n/a	
NIQVIT_1o		0.10167	0.99947
NIQVIT_2o		0.3545	0.99358
NIRRAH_ho		0.20923	0.99832
NIVBIO_ho		0.27073	0.99835
NOBDUFo		0.23058	0.99871
NOKGAXo		0.09112	0.99906
NORGES01o		0.02757	0.99994
NUJDAX_wo		0.10144	0.9988
NUPQEWo		0.08552	0.9997
NUYHUKo		0.04418	0.99988
OBEQAN01o		0.06784	0.9997
OCAHACo		0.06806	0.99971
OFAZUQo		0.17568	0.99792
OFEJOAo		0.03568	0.99989
OFILWVo		0.05666	0.99982
OGIXINo		0.01673	0.99991
OPOGADo		0.02419	0.99995
OPOQERo		0.02804	0.99992
ORNHCL12o		0.06706	0.99916
OROZAXo		0.08453	0.99938

OXYTET01_ho		0.12604	0.99881
PABHIJ01o		0.06994	0.9994
PAJNUJo		0.04815	0.9998
PANCURo		0.11263	0.99954
PBUMSLo		0.20896	0.99732
PBUMSL_ho		0.20896	0.99732
PCLPMS_1o		0.28432	0.99551
PCLPMS_2o		0.16634	0.99848
PEBGUC02o		0.105	0.99895
PEDWOMo		0.109	0.99939
PEKFANo		0.10696	0.99897
PERPAZo		0.08545	0.9996
PETHIDo		0.06759	0.99946
PETHID_ho		0.06759	0.99946
PEXKOS01o	0.0552 wo Cl-ion, OK		n/a
PEYGOQo		0.11195	0.99883
PGDEDOo		0.03546	0.9999
PHEPHRo		0.05242	0.99962
PICVUU01o		0.10304	0.99959
PIGSOOo		0.0394	0.99984
PILOCP01o		0.21653	0.9943
PIMKOLo		0.05709	0.99963
PIMOZDo		0.19124	0.99879
PINDONo		0.05678	0.99963
PIPAMPo		0.13724	0.99935
PIPCIL_ho		0.55519	0.98834
PIRENA10_wo		0.07397	0.99962
PIWLEM02o		0.04117	0.9999
PMEPEN01o		0.06593	0.99971
POBKOG0o		0.11535	0.99945
POGKOMo		0.17417	0.99936
PRAZAM_ho		0.10019	0.99901
PRGDOL02o		0.04603	0.99985
PROCPHo		0.10914	0.99901
PROGLE20_ho		0.36106	0.9923
PROGST12o		0.05848	0.99974
PUFGU0o		0.10077	0.99899
PUQGUFo	0.0778 wo K-ion, OK		n/a
PUVMAUo		0.06904	0.99902
PUVRIHo		0.05053	0.99974
PYRXCL01o		0.0443	0.99964
PYRZIN23o		0.02084	0.99989
QABCIIo		0.13095	0.9995
QABTOD01o		0.04561	0.9999
QATNEF01o		0.09152	0.99963
QECHOV03o		0.02692	0.99981
QEMLIEo		0.08323	0.99964
QETZULo		0.16113	0.99813
QICMEU_ho		1.3643	0.91064
QIFKEX01o		0.28505	0.99635

QIJPACo	0.11878	0.99904
QIJZOYo	0.07371	0.99977
QIQKEIo	0.06937	0.99966
QIQLILo	0.14224	0.99728
QOMKUAo	0.13853	0.99918
QQQAEJ02o	0.12456	0.99753
QQQAUG04o	0.17424	0.99649
QQQBSS01_ho	0.18443	0.9971
QUBQIO01o	0.04653	0.99979
QURWOQ02o	0.07866	0.99916
QUZWOYo	0.14467	0.99846
RALMIAo	0.11174	0.99918
RALMOG01o	0.09553	0.99944
RAVPUBo	0.11706	0.99963
RBFLAV10_ho	0.05375	0.99979
REBCILo	0.06858	0.99984
REDNIXo	0.02118	0.99997
REZBILo	0.07094	0.99952
RIGDESo	0.05105	0.9997
RILQOUo	0.12598	0.99853
RINTITo	0.05982	0.99986
RIQFOOo	0.02523	0.99988
RIWKOXo	0.05359	0.99966
ROKNUBo	0.115	0.99955
ROQLOAo	0.02756	0.99994
RSERPNO1o	0.06414	0.99988
RUKBOPo	0.04324	0.9998
RUKQOFo	0.06983	0.99977
RUTGOEo	0.10642	0.99846
RUTGUKo	0.13208	0.99574
RUWBAMo	0.11541	0.99934
RUYZAOo	0.14751	0.99765
SAFNIWo	0.18995	0.99778
SAHSOJ_ho	0.0844	0.9995
SAJFITo	0.07058	0.9997
SAJNOG_ho	0.12709	0.99927
SAQYIRo	0.11339	0.99948
SATHOLo	0.08193	0.99944
SAWFIF_ho	0.15208	0.99892
SAXFEDo	0.1566	0.99931
SAXQITo	0.09258	0.99911
SAZLAGo	0.07774	0.99948
SAZNOW_mo	1.04061	0.94459
SEBPEU01o	0.07161	0.99924
SENJIFo	0.11156	0.99932
SEQKAA02o	0.08854	0.99922
SEQNILo	0.08365	0.99974
SESKUY01o	0.10766	0.99919
SETSOAo	0.10801	0.99885
SFDMOX05_ho	0.07209	0.99954

SIFGODo		0.04594	0.99988
SIJWUFo		0.02562	0.99982
SIKLIH07o		0.06758	0.99951
SIWCEG_1o		0.58071	0.98471
SIWCEG_2o	2 nd disorder component, OK		n/a
SLFNMB02o		0.07372	0.99938
SLFNMC20_ho		0.4365	0.98495
SLFNMG01o		0.05218	0.99961
SOBBUG_ho		0.09966	0.99864
SONKEMo		0.07659	0.99973
SOPSEVo		0.12329	0.99735
SOYMID_1o		0.12783	0.99931
SOYMID_2o		0.12481	0.99934
SUTHAZ37o		0.09876	0.99887
SUTWIOo		0.16617	0.99835
SUVGULo		0.07814	0.99964
SUWNUTo		0.04281	0.99978
TADLIU02o		0.06383	0.99979
TAJFOZ_ho		0.15295	0.99876
TANDETo		0.06101	0.9991
TAYFUV_ho		0.17324	0.99783
TAZZUPo		0.06631	0.99955
TEPSAM_ho		0.05485	0.9994
TESTON10o		0.01998	0.99996
TETCYH12o		0.65366	0.97227
TEZZEEo		0.05745	0.99973
TFPROM10o		0.11861	0.99898
THCLCS_ho		0.27221	0.99678
THGUAN10o		0.03158	0.99978
TIJMIIo		0.08151	0.99919
TIPWAQ_ho		0.29034	0.99419
TITHYN10_ho		0.09802	0.99944
TIYQAU01o		0.08499	0.99966
TMPHCL01o		0.09478	0.99928
TOPXED_mo		0.11378	0.99893
TOXXIN_ho		0.2012	0.99729
TOYJOGo		0.16045	0.99874
TOYSUXo		0.30231	0.99469
TUBOBM_ho		0.30468	0.9957
TULRIB_mo	0.0138 for main molecule, OK		n/a
TUQGETo		0.30116	0.98274
TUTDODo		0.06265	0.99971
TUZCOIo		0.05442	0.99976
UCUVETo		0.07066	0.9996
UGIYEPo		0.07854	0.99971
UHUQUKo		0.0986	0.9997
UKAKIAo		0.10855	0.99912
UMOMALo		0.06128	0.99982
UPUKOHo		0.08006	0.99923
UQAPEJo		0.09666	0.9995

URAVUGo		0.11861	0.99939
UREAOH12o		0.04305	0.99903
USOCOVo		0.12698	0.99947
UXIXUVo		0.04642	0.99966
VAMBOA01o		0.1859	0.99881
VAWPITo		0.102	0.99851
VAWQEQo		0.07488	0.99957
VAYJOWo		0.2376	0.99689
VAYXOIo		0.1061	0.99945
VEMRAIo		0.10869	0.99951
VEVQAQo		0.08667	0.99955
VICXOVo		0.1236	0.99919
VIDJEX01o		0.15361	0.99784
VIFQILo		0.10361	0.99928
VIGDAR_1o		0.22684	0.99679
VIGDAR_2o		0.23294	0.99662
VOGCAV_ho		0.08841	0.99982
VOXDUHo		0.13282	0.99819
VUCDUSo		0.24421	0.99307
VUCJElO		0.11824	0.99928
VUJRUE_ho		0.05639	0.99989
VUXPUZo		0.09748	0.99861
WACHIR_ho		0.08044	0.99977
WADPICo		0.0886	0.99954
WADTIGo		0.06208	0.99981
WAHJARo		0.10273	0.99955
WAJYUCo		0.11277	0.99959
WALPIJo		0.10863	0.99899
WASGAA_ho		0.08781	0.99945
WAWVOHo		0.10572	0.99955
WEHYEOo		0.08654	0.99942
WELGAVo		0.08137	0.99968
WEMGECo		0.05612	0.99951
WEQHEF01_ho		0.67036	0.96789
WERVEU02o		0.093	0.99923
WETLOYo		0.08313	0.99949
WETPOB_1o	0.054, disordered, OK		n/a
WETPOB_2o	0.082, disordered, OK		n/a
WEXQAS01o		0.07044	0.99958
WEZCOTo		0.07175	0.99956
WIBZEMo		0.1187	0.99944
WINJAE_mo	0.0284, shifted due to cation, OK		n/a
WINWUL02o		0.08977	0.99933
WIPYOLo		0.10867	0.99904
WOQDAHo		0.07312	0.99947
WOYPABo		0.07603	0.99967
WUTHEA01o	Moved molecules in unit cell, higher Z structure, OK		n/a
XANTOX_ho		0.18893	0.99512
XAPTAK01o		0.03314	0.99986
XAQNOUo		0.09623	0.99885

XATJAFo		0.08216	0.99953
XAYGEJ01o		0.11048	0.99911
XAZRUNo		0.10032	0.99901
XEJKUSo		0.07451	0.99888
XEZTON01o		0.09007	0.99943
XICRUYo		0.12032	0.99851
XIJKIK01o		0.04537	0.99898
XIRHOXo		0.13245	0.99699
XOCXUI01o		0.06183	0.99972
XOGMOXo		0.14906	0.99565
XOJNOBo		0.13887	0.99889
XOSGUHo		0.07944	0.99978
XOZRUZo		0.24582	0.99525
XULRUTo		0.06772	0.99975
YACGUE_ho		0.32192	0.99121
YACHAL_ho		0.09409	0.99937
YACTEC01o		0.06532	0.99944
YALWISo		0.04813	0.99989
YALYUHo		0.0736	0.99963
YAPZEUo		0.03839	0.99989
YARXEWo		0.05939	0.99986
YIGRAIo		0.06347	0.99961
YIKNUDo		0.11619	0.99887
YODXARo		0.13084	0.99879
YOKQAQo		0.0937	0.99852
YUKVADo		0.04393	0.99989
YURSUB_ho		0.12828	0.99937
ZAPLEJo		0.11601	0.9995
ZAPXOFo		0.16955	0.99574
ZEMJOQo		0.05147	0.99984
ZENREPo		0.15334	0.99939
ZIDLEDo		0.07708	0.99964
ZIKJOSo		0.0599	0.99949
ZIMBOOo		0.077	0.99857
ZIVKEUo		0.09763	0.99907
ZOGSODo		0.22119	0.99727
ZOXSEKo		0.1117	0.99914
ZUHFIRo		0.08305	0.99935
ZUQMIJ_ho	0.105, OK		n/a
ZZPUS19o		0.14205	0.99801
ZZZRCG01_ho		0.09679	0.99942
ZZZTSE03o		0.04035	0.9998























































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































































