Supplementary information for

β -Cyclodextrin dimethylformamide 12.5 hydrate: a deeper insight into β -cyclodextrin crystal packing

Rubén Granero-García and Francesca P. A. Fabbiani*

Table of contents

S1	Structure coordinates of β -CD·DMF 12.5 from invariom refinement	1
S2	Structures used in the XPAC calculation	5
S3	Hydrogen-bond distances and angles in the structure of β -CD·DMF 12.5 hydrate	6
S4	Results of the CLPDIM calculation	7
S5	Complete data from the XPAC calculation	9

Section S1 Structure coordinates of β -CD·DMF 12.5 hydrate from invariom refinement used in the CLPCRY calculation.

Page

loop_ _atom_site_label _atom_site_type_symbol _atom_site_fract_x _atom_site_fract_y _atom_site_fract_z 03 0 0.2767 0.3385 0.33916 05 0 0.1718 0.4715 0.24995 07 0 0.1947 0.76044 0.24738 09 0 0.2509 0.6770 0.40630 014 0 0.1073 0.8531 0.3949 018 0 0.6430 0.3534 0.33098 020 0 0.4561 0.3296 0.32944 022 0 0.3718 0.5737 0.36465 024 0 0.5774 0.6032 0.44511 027 0 0.4210 0.6493 0.51610 030 0 0.9374 0.6477 0.26766 032 0 0.8035 0.4893 0.31979 034 0 0.6513 0.6288 0.35346 036 0 0.8100 0.9034 0.34674 039 0 0.7108 0.8924 0.45731 042 0 0.8620 1.0507 0.03610 044 0 0.9145 0.8479 0.11958 046 0 0.8341 0.8755 0.24075 048 0 0.8499 1.2302 0.18235 051 0 0.9437 1.1844 0.29310 054 0 0.5431 1.2074 -0.05783 056 0 0.7125 1.1335 -0.01483 058 0 0.7409 1.1791 0.11313 060 0 0.5719 1.4274 0.07600 063 0 0.7133 1.5365 0.14223

066 0 0.1913 1.1010 -0.01438 068 0 0.3578 1.2145 -0.05237 070 0 0.4672 1.2667 0.05179 072 0 0.2512 1.3204 0.11512 075 0 0.3552 1.5508 0.13592 078 0 0.1548 0.6235 0.14052 080 0 0.1842 0.8449 0.05324 082 0 0.2307 1.0887 0.10862 084 0 0.0943 0.9302 0.22747 087 0 0.0539 1.2100 0.19257 C1 C 0.2915 0.5515 0.3948 C2 C 0.2359 0.4663 0.3503 C4 C 0.2266 0.5433 0.2908 C6 C 0.1894 0.6857 0.3033 C8 C 0.2394 0.7621 0.3531 C10 C 0.1923 0.8870 0.3743 C16 C 0.6307 0.5386 0.4014 C17 C 0.5867 0.4116 0.3744 C19 C 0.4995 0.4500 0.3477 C21 C 0.4479 0.5282 0.3947 C23 C 0.4980 0.6510 0.4186 C25 C 0.4518 0.7329 0.4675 C28 C 0.8659 0.8533 0.3002 C29 C 0.8783 0.7000 0.3105 C31 C 0.7913 0.6286 0.3079 C33 C 0.7339 0.6922 0.3570 C35 C 0.7252 0.8452 0.3470 C37 C 0.6757 0.9190 0.3972 C40 C 0.8300 1.2040 0.1212 C41 C 0.8806 1.0764 0.0989 C43 C 0.8582 0.9531 0.1389 C45 C 0.8705 0.9886 0.2066 C47 C 0.8267 1.1253 0.2248 C49 C 0.8511 1.1790 0.2874 C52 C 0.5276 1.3585 0.0284 C53 C 0.5917 1.2823 -0.0127 C55 C 0.6498 1.1907 0.0259 C57 C 0.6921 1.2722 0.0774 C59 C 0.6222 1.3421 0.1159 C61 C 0.6582 1.4347 0.1654 C64 C 0.2078 1.2136 0.0827 C65 C 0.2345 1.2098 0.0146 C67 C 0.3331 1.1975 0.0100 C69 C 0.3780 1.3034 0.0512 C71 C 0.3426 1.3037 0.1161 C73 C 0.3775 1.4198 0.1577 C76 C 0.1162 0.7911 0.2175 C77 C 0.1248 0.7595 0.1491 C79 C 0.1873 0.8593 0.1187 C81 C 0.1628 1.0034 0.1341 C83 C 0.1572 1.0215 0.2040 C85 C 0.1312 1.1655 0.2226 H1 H 0.3020 0.4956 0.4382 H2 H 0.1719 0.4492 0.3706

H3 H 0.2824 0.2887 0.37711 H4 H 0.2905 0.5544 0.2697 H5 H 0.1660 0.5213 0.21202 H6 H 0.1215 0.6762 0.3171 H8 H 0.3029 0.7917 0.3351 H10 H 0.2281 0.9344 0.4124 H11 H 0.1883 0.9612 0.3365 H14 H 0.1101 0.8158 0.4358 H16 H 0.6910 0.5071 0.4241 H17 H 0.5766 0.3377 0.4118 H18 H 0.6214 0.2659 0.31827 H19 H 0.5096 0.5146 0.3069 H20 H 0.4009 0.3525 0.31205 H21 H 0.4309 0.4610 0.4334 H23 H 0.5129 0.7182 0.3795 H25 H 0.4961 0.8095 0.4864 H26 H 0.3973 0.7869 0.4463 H27 H 0.4686 0.6037 0.53558 H28 H 0.9289 0.9040 0.3047 H29 H 0.9051 0.6844 0.3569 H30 H 0.9108 0.6429 0.22765 H31 H 0.7618 0.6425 0.2622 H32 H 0.7498 0.4414 0.31476 H33 H 0.7620 0.6730 0.4028 H35 H 0.6938 0.8642 0.3023 H37 H 0.6772 1.0292 0.3884 H38 H 0.6081 0.8860 0.3961 H39 H 0.7697 0.9229 0.45879 H40 H 0.8485 1.2921 0.0927 H41 H 0.9499 1.0971 0.1035 H42 H 0.9149 1.0468 0.01316 H43 H 0.7911 0.9225 0.1304 H44 H 0.8930 0.7612 0.13354 H45 H 0.9398 0.9956 0.2167 H47 H 0.7564 1.1116 0.2230 H49 H 0.8239 1.2814 0.2937 H50 H 0.8246 1.1122 0.3232 H51 H 0.9613 1.2766 0.30178 H52 H 0.4932 1.4341 0.0000 H53 H 0.6322 1.3573 -0.0367 H54 H 0.5819 1.1582 -0.08425 H55 H 0.6113 1.1080 0.0462 H56 H 0.7611 1.1011 0.00855 H57 H 0.7349 1.3496 0.0574 H59 H 0.5805 1.2642 0.1368 H61 H 0.6045 1.4834 0.1898 H62 H 0.6939 1.3724 0.1989 H63 H 0.6819 1.5933 0.11386 H64 H 0.1375 1.2278 0.0859 H65 H 0.2146 1.3059 -0.0073 H66 H 0.1925 1.0217 0.01169 H67 H 0.3524 1.0951 0.0255 H68 H 0.4196 1.2188 -0.05519 H69 H 0.3708 1.4054 0.0307

H71 H 0.3578 1.2052 0.1378 H73 H 0.3511 1.4068 0.2042 H74 H 0.4479 1.4114 0.1609 H75 H 0.2947 1.5541 0.12645 H76 H 0.0652 0.7261 0.2371 H77 H 0.0612 0.7696 0.1272 H78 H 0.1094 0.5685 0.12332 H79 H 0.2531 0.8385 0.1351 H80 H 0.2213 0.7714 0.04078 H81 H 0.1009 1.0297 0.1123 H83 H 0.2202 0.9977 0.2246 H85 H 0.1211 1.1674 0.2726 H86 H 0.1837 1.2366 0.2119 H87 H 0.0061 1.2034 0.22055 O1A O 0.603145 1.133141 0.261041 N1A N 0.53263 0.951339 0.222338 C2A C 0.542102 1.06064 0.258549 C3A C 0.595147 0.912863 0.178293 C4A C 0.458994 0.860751 0.225827 H2A H 0.487602 1.085903 0.289204 H3A H 0.639487 0.998402 0.170333 H3B H 0.562939 0.885758 0.134865 H3C H 0.631559 0.824552 0.195481 H4A H 0.439444 0.84906 0.274143 H4B H 0.476419 0.760611 0.206864 H4C H 0.405295 0.903905 0.199119 0111 0 1.018051 1.428764 0.311622 H112 H 1.072124 1.460853 0.295493 H113 H 0.971226 1.482951 0.295797 0211 0 0.883074 0.586837 0.1554 H212 H 0.915252 0.503618 0.150421 H213 H 0.823199 0.571994 0.145777 0311 0 0.279354 1.19584 0.45014 H312 H 0.29206 1.273336 0.475838 H313 H 0.222185 1.163762 0.459393 0411 0 0.55565 0.593391 0.59352 H412 H 0.56342 0.67542 0.61727 H413 H 0.610196 0.561461 0.578154 0511 0 -0.004529 1.061733 0.397638 H512 H -0.013244 1.121587 0.363024 H513 H 0.053078 1.026043 0.394291 0611 0 0.024476 1.439739 0.122711 H612 H 0.032646 1.357201 0.146029 H613 H 0.005893 1.418169 0.081506 0711 0 0.107977 0.775609 0.512152 H712 H 0.086156 0.866547 0.518183 H713 H 0.059086 0.719739 0.501621

	<i>a</i> /Å	<i>b</i> /Å	c /Å β /°		V /Å ³	Guest	Hydration number				
ARUXIU	15.1711(3)	10.16870(10)	20.9862(4)	110.92	3024.13	formic acid	7.7				
ARUXOA	15.263(4)	10.157(2)	21.044(5)	110.698(4)	3051.81	acetic acid	7.7				
BCDEXD03	21.283(5)	10.322(1)	15.092(5)	112.41(1)	3065.07		12.26				
BCDEXD04	21.161(7)	10.254(1)	15.110(7)	111.91(2)	3041.83		11.56				
BCDEXD05	21.233(5)	10.294(1)	15.103(4)	112.22(1)	3055.96		11.89				
BCDEXD10	21.29(2)	10.33(1)	15.10(2)	112.3(5)	3072.51		12				
BOBPEN	21.25(3)	10.28(2)	15.30(2)	113.25(9)	3070.86	hydrogen iodide	8				
BOBPIR	21.03(3)	10.11(2)	15.33(2)	111.02(9)	3042.47	methanol	6.5				
BUVSEQ01	21.080(7)	10.197(1)	15.131(6)	111.58(2)	3024.47		11.19				
DEBGOG	15.356(3)	10.101(2)	21.287(3)	112.81(1)	3043.63	benzyl alcohol	5				
DEWCOY	15.4244(16)	10.1574(11)	20.557(2)	110.074(2)	3025.04	benzamide	6				
DIRVOP	15.285(3)	10.345(2)	20.118(2)	102.14(1)	3109.98	hexamethylenetetramine	6				
GUXZOO	15.1220(4)	10.3350(2)	20.9150(3)	109.575(3)	3079.8	<i>m</i> -aminophenol	7.5				
HAQNUK	15.274(12)	10.178(8)	20.961(16)	110.251(7)	3057.14	1,4-dioxane	7				
IZALEB	15.0678(9)	10.2126(6)	20.9653(13)	111.7473(12)	2996.55	nicotinic acid	7.65				
KIJSEC	15.1977(7)	10.1763(5)	20.6943(6)	109.239(4)	3021.76	succinic acid	7				
KOBRIC	15.223(5)	10.578(3)	20.204(6)	108.37(7)	3087.64	potassium hydroxide	8				
KUTKOZ	21.199(12)	9.973(3)	15.271(8)	110.87(3)	3016.74	1,4-butanediol	6.25				
LADBAV	15.15(2)	10.343(15)	20.93(3)	109.69(4)	3087.89	pentanedioic acid	4.2				
LADQOX	20.82(1)	10.366(6)	14.968(9)	110.12(4)	3033.26	diaqua-dichloro-copper(II)	8.9				
MIFHAK	15.231(3)	10.068(3)	21.117(3)	110.16(5)	3039.81	squaric acid	6.65				
NUFVEQ	20.8314(9)	10.2937(6)	15.2155(8)	110.733(2)	3051.4	pyrazinamide	5.5				
NUFVIU	20.95040(10)	10.2493(2)	15.1424(3)	108.7630(10)	3078.69	piperazine	7.2				
PIJGIY	21.212(1)	10.021(3)	15.208(1)	111.47(3)	3008.38	ethylene glycol	8				
PIJGOE	21.322(6)	9.954(1)	15.251(6)	111.20(1)	3017.8	glycerol	7.2				
POBRON	20.857(6)	10.158(1)	15.140(6)	110.94(1)	2995.79		9.35				
POBRON01	15.107(7)	10.242(3)	20.911(8)	110.27(3)	3035.1		9				
POVSIC	20.042(6)	10.378(2)	15.123(9)	102.30(2)	3073.32	trans-cyclohexane-1,4-diol	5.4				
SUJDAD	20.8353(4)	9.93970(10)	15.2043(3)	110.630(2)	2946.85		10.41				
TUSHUL	15.1667(5)	10.1850(3)	20.9694(7)	110.993(2)	3024.2		7.5				
UPULEX	20.7890	10.2084	15.1091	110.8250	2997.01	4-aminobenzoic acid	10				
VACZIJ	15.1550(3)	10.2850(2)	20.9056(5)	109.86	3064.74	dimethylsulfoxide	7.35				
VOJLIQ	15.2196(10)	10.2279(7)	20.9321(14)	110.9270(10)	3043.45	p-aminobenzoic acid	5				
XEGTAF	10.601(4)	26.402(9)	15.586(6)	93.145(6)	4355.76	6'-Methoxycinchonan-9-ol	16.5				
YIYSII	21.310	9.987	15.247	111.85	3011.8	diethanolamine	6.4				
YIYSOO	21.451	10.014	15.240	111.25	3051.12	1,5-pentanediol	6.2				
ZIGZIY	20.988(9)	10.092(2)	15.223(10)	110.30(6)	3024.13	but-2-yne-1,4-diol	7				
β-CD·DMF	15.500(3)	9.794(2)	21.689(5)	90.138(3)	3292.5(12)	dimethylformamide	12.5				

Section S2 $P2_1$ structures used in the XPAC calculation with unit-cell parameters, guest and number of water molecules of hydration. Structures highlighted in grey are those in which the *a* axis is longer than the *c* axis. In all the cases the *b* axis is the shortest axis, with the exception of the refcode highlighted in yellow.

H bond	D-H	H···A	D ···A	<i>D</i> -H··· <i>A</i>
O2_1—H2O_1O3_9 ^{1_545}	0.84	1.95	2.785(6)	173.5
O3_1—H3O_1O2_7	0.84	1.98	2.816(5)	177.7
O6_1—H6O_1O7_9	0.84	1.81	2.643(11)	170.8
O6' 1—H6O' 1O8' 9 ^{2_646}	0.84	2.10	2.92(6)	169
O2 2—H2O 2O1 $8^{1}-545$	0.84	1.92	2.707(6)	154.4
O3 2—H3O 2O2 1	0.84	2.10	2.791(5)	138.6
O6 2—H6O 2O4 9	0.84	1.95	2.733(6)	154.0
O2 3—H2O 3O2 9	0.84	1.82	2.638(5)	162.5
O3 3—H3O 3…O2 2	0.84	2.01	2.828(5)	163.9
O6 3—H6O 3O9 9	0.84	1.93	2.743(9)	161.6
O6 3—H6O 3O9' 9	0.84	1.97	2.784(18)	162.6
O2 4—H2O 4O10 9	0.84	2.03	2.711(9)	137.7
O2 4—H2O 4O10' 9	0.84	1.90	2.691(8)	155.6
O3 4—H3O 4O2 9	0.84	1.90	2.723(5)	165.3
O6 4—H6O 4O1 9	0.84	1.86	2.688(6)	166.9
O2 5—H2O 5O6 6 ^{2_645}	0.84	1.94	2.776(5)	173.6
O3 5—H3O 5O2 4	0.84	1.86	2.689(5)	168.3
O6 5—H6O 5O3 6 ² -6 ⁵⁵	0.84	2.01	2.835(5)	169.4
O2 6—H2O 6O3 7	0.84	2.08	2.916(5)	171.1
O3 6—H3O 6O2 5	0.84	2.04	2.875(5)	174.2
$06^{-}6-H60^{-}602^{-}7^{1-565}$	0.84	2.40	3.188(5)	156.0
O2 7—H2O 7O6 9 ^{1_545}	0.84	1.94	2.733(6)	158.2
O3 7—H3O 7O3 5 ² _645	0.84	1.90	2.743(5)	177.9
$067 - H607 \dots 064^{1}$	0.84	1.98	2.786(5)	161.9
01^{9} -H1A 9 03^{1} -1 ⁻⁶⁶⁵	0.84(3)	1.93(3)	2.766(5)	173(6)
O1 9—H1B 9O2 3 ^{1_565}	0.85(3)	1.87(4)	2.652(5)	151(6)
O2 9—H2A 9O6 9 ^{1_645}	0.86(3)	1.96(4)	2.721(6)	147(6)
O2 9—H2B 9O6 5^{1}	0.87(3)	1.86(4)	2.686(5)	157(6)
O3 9—H3A 9O6 3 ² _ ⁶⁵⁶	0.91(3)	1.90(4)	2.776(6)	161(6)
O3 9—H3B 9O11' 9	0.90(3)	2.04(4)	2.93(2)	170(7)
O3 9—H3B 9O11" 9	0.90(3)	1.90(4)	2.770(9)	162(7)
04^{9} -H4A 9 03^{2}_{-656}	0.86(3)	2.03(3)	2.842(6)	159(7)
O5 9—H5A 9O6 4 ^{1_455}	0.89(3)	1.84(4)	2.685(6)	156(7)
O5 9—H5B 9O6 1	0.86(3)	1.96(5)	2.676(7)	139(7)
O6 9—H6A 9O6 7	0.88(3)	1.87(3)	2.748(6)	176(7)
$06^{-}9-H6B^{-}9010'^{-}9^{2}_{-}655$	0.88(3)	1.73(4)	2.549(9)	153(7)
06^{9} -H6B 9010" 9^{2} -655	0.88(3)	2.09(4)	2.931(9)	159(7)
O7 9—H7A 9O12 9 ^{2_556}	0.87(3)	2.05(13)	2.552(19)	116(11)
O7 9—H7A 9O12' 9 ^{2_556}	0.87(3)	1.91(7)	2.739(18)	159(14)
O7 9—H7B 9O12" 9	0.86(3)	2.39(16)	2.832(18)	112(13)
08 9—H8A 906 2 ² ₋₆₅₆	0.85(7)	2.10(8)	2.819(13)	143(7)
O8 9—H8B 9O2 2 ^{1_565}	0.87(11)	2.17(11)	3.007(12)	163(11)
O8' 9—H8A 9O6 2 ^{2_656}	0.91(7)	2.10(8)	2.78(2)	131(6)
O8'_9—H8B'_9O7 9 ^{2_656}	0.87(6)	1.63(10)	2.30(2)	131(16)
O9_9—H9A_9O5_9 ^{1_655}	0.85(5)	2.01(5)	2.832(10)	162(5)
O9_9—H9B_9O11" 9 ^{2_646}	0.86(3)	2.07(7)	2.829(13)	147(10)
O9'_9—H9A_9O5_9 ^{1_655}	0.89(6)	2.01(5)	2.762(18)	141(5)
O9'_9—H9B'_9O8_9	0.89(14)	2.31(18)	3.14(2)	157(17)

Section S3 Hydrogen-bond distances (in Å) and angles (in °) in the structure of β -CD·DMF 12.5 hydrate.

Section S4 Results of the CLPDIM calculation.



Figure S4.1 Cartesian reference system used in the CLPDIM calculation.



Figure S4.2 Total host-guest interaction energies for translations (along the *x*-, *y*- and *z*-axes) of the DMF with respect to the β -CD cavity, as illustrated in Figure S4.1. The point of zero translation corresponds to the the host-guest geometry observed experimentally.



Figure S4.3 Total host-guest interaction energies for the combination of translations and rotations (along and about the *x*-, *y*- and *z*-axes) of the DMF with respect to the β -CD. Energy values are colour coded.

Section S5	Complete data	from the XPA	C calculation.																																			
	ARUXIU	ARUXOA	BCDEXD03	BCDEXD04	BCDEXD05	BCDEXD10	BOBPEN	BUVSEQ01	DEBGOG	DEWCOY	DIRVOP	GUXZOO	HAQNUK	IZALEB	KIJSEC	KOBRIC	KUTKOZ	LADBAV	LADQOX	MIFHAK	NUFVEQ	NUFVIU	PIJGIY	PIJGOE	POBRON	POBRON01	POVSIC	SUJDAD	TUSHUL	UPULEX	VACZIJ	NOJLIQ	XEGTAF	ЛY	VIYSOO	ZIGZIY	β-CD·DMF	
ARUXIU																																						ARUXOA
ARUXOA	3D 0.6 0.1 14 21																																					ARUXOA
BCDEXD03	0.3 0.6 3D 0.1 1.3 0.1 12 21	3D 1.5 0.1 12 21																																				BCDEXD03
BCDEXD04	0.7 1.1 3D 0.8 0.1 12 21 21	0.7 1.3 3D 1.1 0.1 12 21	3D 0.6 0.1 12 21																																			BCDEXD04
BCDEXD05	0.4 0.7 3D 1.1 0.1 12 21	0.5 1 3D 1.3 0.1 12 21	0.3 0.5 3D 0.4 0 12 21	3D 0.3 0 12 21									Parame	ters for each	comparison:								г															BCDEXD05
BCDEXD10	0.5 1 3D 1.6 0.1 12 21	0.6 1.1 3D 1.7 0.1 12 21	0.2 0.3 3D 0.6 0 12 21	0.2 0.3 3D 0.9 0 12 21	3D 0.7 0 12 21								SCA X n δ(a) δ	N D p D(p)	SCA N X D D S n N	umber of dim issimilarity ind tretch parame umber of neig	nensions in w dex eter (in Å) ghbours	hich there is a	a similarity (N	No, 1D, 2D o	r 3D)		b															BCDEXD10
BOBPEN	0.7 1.4 3D 3.1 0.1 14 21	0.8 1.5 3D 3 0.1 14 21	0.2 0.6 3D 3.8 0.1 12 21	0.4 0.8 3D 3.5 0.1 12 21	0.2 0.6 3D 3.7 0.1 12 21	3D 4 0.1 12 21									p Ν δ(a) Μ δ(a) Μ	umber of poir lean differenc lean differenc	nts per mole ce between a ce between i	cule angles (in °) nterplanar ang	gles (in °)				Г															BOBPEN
BOBPIR	1.3 2.9 3D 1 1 0.1 14 21	1.2 2.7 3D 1 0.1 14 21	1.5 3.5 3D 2.1 0.2 12 21	1.4 3.2 3D 1.6 0.2 12 21	1.4 3.4 3D 1.9 0.2 12 21	1.5 3.6 3D 3D 2.3 0.2 3 12 21 1	3D 3 0.2 14 21																															BOBPIR
BUVSEQ01	0.5 0.9 3D 0.6 0 14 21 21	0.4 0.9 3D 0.9 0.1 14 21	1.1 1.8 3D 0.9 0.1 12 21	0.8 1.4 3D 0.4 0 12 21	0.9 1.6 3D 0.7 0.1 12 21	1.1 2 1 3D 3D 1.2 0.1 3 12 21 1	.3 2.7 3D 3E 5.4 0.1 1.4 14 21 14	0.1 21															г															BUVSEQ01
DEBGOG	0.2 0.5 3D 0.2 2.1 0.2 14 21	0.4 0.8 3D 2.1 0.1 14 21	0.5 0.8 3D 3.3 0.3 12 21	0.2 0.4 3D 0.3 12 21	0.4 0.6 3D 0.3 12 21	0.5 1.1 1 3D 3 3 12 21 1	.3 3.1 0.7 3D 3E 5.5 0.2 1.5 14 21 14	1.2 3D 0.1 2.5 21 14).2 21																													DEBGOG
DEWCOY	1 1.9 No 	1 1.9 No 	1.6 2.8 No 	1.4 2.5 No 	1.5 2.7 No 	1.7 3 1 No 	.6 3 0.7 No No 	1.3 1.2 2 No 	2.2 														Г															DEWCOY
DIRVOP	1D 7.2 0.2 2 21	1D 7.4 0.2 2 21	1D 7 0 2 21	1D 7.1 0.1 2 21	1D 7.1 0.1 2 21	1D 7.3 0 2 21	1D 1E 7.8 0.1 7.7 2 21 2	1D 0.2 7.1 0 21 2 2	1D 0.1 7.8 0.2 21 2 21	No 2 																												DIRVOP
GUXZOO	4.5 5.6 3D 1.3 0.1 12 21	4.7 5.8 3D 1.4 0.1 12 21	4.4 5.5 3D 0.4 0 12 21	4.4 5.5 3D 0.7 0.1 12 21	4.4 5.5 3D 0.5 0 12 21	4.5 5.7 4 3D 0.7 0 3 12 21 1	.9 6.1 4.7 3D 3E .8 0.1 2.1 12 21 12	6.1 4.4 5 0 3D 0.2 1 0 21 12 2	5.6 4.8 6.2 3D 0.1 3.3 0.3 21 12 21	2 No 3 1	1D 7.1 0 2 21						T						Г															GUXZOO
HAQNUK	0.7 1.1 3D 1.2 0.1 14 21	0.7 1.3 3D 1 0 14 21	0.1 0.4 3D 1.2 0.1 12 21	0.3 0.6 3D 1.1 0.1 12 21	0.2 0.4 3D 1.2 0.1 12 21	0.2 0.6 1 3D 1.5 0.1 3 12 21 1	.5 3.5 1.1 3D 3E 3.3 0.1 1.7 4 21 14	1.8 0.5 0 3D 3D 0.1 1.1 0 21 14 2 2 14 2	0.8 1.6 2.8 3D 0.1 2.9 0.2 21 14 21	3 No 2 1	4.4 5.6 1D 7.3 0.2 2 21	3D 1.3 0.1 12 21																										HAQNUK
IZALEB	0.5 1.1 3D 1 0.1 14 21	0.4 0.9 3D 1.2 0.1 14 21	0.6 1.1 3D 0.8 0.1 12 21	0.5 1 3D 0.6 0.1 12 21	0.5 1 3D 0.6 0.1 12 21 21	0.7 1.3 1 3D 1.1 0.1 3 12 21 1	.3 3 0.7 3D 3E 5.6 0.1 1.8 14 21 14	1.6 0.5 3D 3D 0.2 0.7 0 21 14 2	1 1.3 2.6 3D 0.1 2.9 0.3 21 14 21	6 No 3 1	4.6 5.7 1D 7.1 0.1 2 21	0.6 1.1 3D 0.8 0.1 12 21	3D 1 0.1 14 21				T						Г															IZALEB
KIJSEC	0.4 0.9 3D 2.5 0.1 14 21	0.5 1.1 3D 2.2 0.1 14 21	0.4 0.7 3D 2.3 0.1 12 21	0.2 0.6 3D 2.4 0.1 12 21	0.3 0.6 3D 2.4 0.1 12 21	0.5 1 1 3D 2.6 0.1 3 12 21 1	.4 3.2 0.8 3D 3E 3.3 0.1 2.9 14 21 14	1.6 0.3 0 3D 3D 3D 0.2 2.5 0 21 14 2	0.6 1.4 2.5 3D 0.1 3.8 0.3 21 14 21	5 No 3 I	4.4 5.5 1D 7.3 0.2 2 21	0.4 0.7 3D 2.4 0.1 12 21	0.5 0.9 3D 1.8 0.1 14 21	3D 2.3 0.1 14 21																								KIJSEC
KOBRIC	1 2.3 No 	0.9 2 No 	1 2.1 No 	1 2.2 No 	0.9 2.2 No 	1 2.3 1 No 	.3 3.1 1.2 No No 	2.6 1 2 • No 	2.3 1.7 3.4 No	4 No 	4.8 5.5 No 	1 2.2 No 	0.7 1.7 No 	0.9 2.1 No 	No 								Г															KOBRIC
KUTKOZ	3D 2 0.1 14 21	3D 1.9 0.1 14 21	3D 2.7 0.2 12 21	3D 2.3 0.2 12 21	3D 2.5 0.2 12 21	3D 2.8 0.2 3 12 21 1	3D 3E 5.7 0.2 1.3 14 21 14	3D 0.1 2.2 0 21 14 2	3D 0.1 2 0.1 21 14 21	No 1 1	1D 8.2 0.4 2 21	3D 2.7 0.2 12 21	3D 2.5 0.1 14 21	3D 2.6 0.2 3 14 21 7	3D 3.6 0.2 14 21	No 																						KUTKOZ
LADBAV	0.8 1.8 3D 1.6 0.1 12 21	0.7 1.8 3D 1.6 0.1 12 21	1.2 2.4 3D 1.1 0 12 21	1 2.1 3D 1.2 0.1 12 21	1.1 2.3 3D 1.1 0 12 21 21	1.2 2.5 1 3D 2 1.2 0 2 12 21 1	.5 3.3 0.5 3D 3E 4 0.1 2.2 12 21 12	1.2 0.9 3D 3D 0.2 1.4 0 21 12 2	2 0.9 1.8 3D 0.1 3.3 0.3 21 12 21	3 No 3 1	4.9 6.6 1D 7.3 0 2 21	1.2 2.4 3D 1 0 12 21	0.9 2.3 3D 1.5 0.1 12 21	1 2.4 1 3D 1.2 0.1 2 12 21 2 2	1.4 3.3 3D 2.6 0.1 12 21	2 2	3D 2.7 0.2 12 21						Г															LADBAV
LADQOX	0.7 1.4 3D 1.8 0.1 12 21	0.7 1.4 3D 2.1 0.2 12 21	0.4 1 3D 1 0.1 12 21	0.5 1.1 3D 1.2 0.1 12 21	0.4 1 3D 1.1 0.1 12 21	0.4 1.2 1 3D 1.3 0.1 4 12 21 1	.5 3.7 1.1 3D 3E .3 0.1 2.6 12 21 12	1.9 0.6 1 0 3D 0.2 1.4 0 21 12 2	1.2 1.6 2.9 3D 0.1 3.6 0.3 21 12 21	9 No 3 1	4.5 5.7 1D 6.7 0 2 21	0.3 0.9 3D 1 0.1 12 21	0.6 1.3 3D 1.9 0.2 12 21	0.5 1.1 3D 1.2 0.1 2 12 21 ⁻	1 2.4 3D 2.9 0.1 12 21	No 3	1.2 2.4 3D 3.3 0.2 12 21	3D 1.4 0.1 12 21																				LADQOX
MIFHAK	0.8 1.6 3D 1.6 0.1 14 21	1 1.8 3D 1.2 0.1 14 21	0.5 0.9 3D 1.8 0.1 12 21	0.5 1.1 3D 1.7 0.1 12 21	0.5 1 3D 1.8 0.1 12 21	0.6 1.1 1 3D 1.9 0.1 3 12 21 1	.7 4 1.2 3D 3E 5.5 0.1 1.9 14 21 14	2.3 0.7 1 0 3D 3D 0.1 1.7 0 21 14 2	1.3 1.8 3.2 3D 0.1 2.9 0.2 21 14 21	2 No 2 1	4.1 5.3 1D 7.8 0.3 2 21	0.5 0.8 3D 1.8 0.1 12 21	0.9 1.7 3D 1.3 0.1 14 21 21	0.6 1.1 1 3D 1.8 0.1 2 14 21	1.3 2.6 3D 2.1 0.1 14 21	No 2 2 	1.4 2.9 3D 2.2 0.1 14 21	0.6 1.3 3D 1.9 0.1 2 12 21	3D 2.5 0.2 12 21				Г															MIFHAK
NUFVEQ	0.7 1.5 3D 1.1 0 12 21	0.5 1.1 3D 1.2 0.1 12 21	0.8 1.6 3D 1 0.1 12 21	0.7 1.5 3D 0.9 0.1 12 21	0.8 1.6 3D 0.9 0.1 12 21	0.9 1.7 1 3D 1.3 0.1 3 12 21 1	.5 3.2 0.7 3D 3E .6 0.1 1.7 12 21 12	1.7 0.7 1 0 3D 0.1 0.9 21 12 2	1.5 1.2 2.6 3D 0 2.8 0.2 21 12 21	6 No 2 1	4.9 6 1D 7.2 0.1 2 21	0.8 1.6 3D 1 0.1 12 21	0.5 1.2 3D 1 0.1 12 21	0.8 1.6 0 3D 0.7 0.1 2 12 21 7	0.9 1.9 3D 2.3 0.1 12 21	(No 2	0.8 2 3D 2.6 0.1 12 21	0.8 1.7 1 3D 1.3 0.1 1 12 21	.1 2.2 3D I.4 0.1 1 12 21	3D 1.8 0.1 12 21																		NUFVEQ
NUFVIU	0.5 1 3D 1.8 0.2 12 21 21	0.5 1.1 3D 1.8 0.2 12 21	0.5 0.8 3D 0.9 0.1 12 21	0.4 0.7 3D 1.2 0.1 12 21	0.4 0.8 3D 1.1 0.1 12 21	0.6 1.1 1 3D 1.2 0.1 3 12 21 1	.3 3.3 0.8 3D 3E .8 0.1 2.5 12 21 12	1.5 0.4 0 3D 3D 0 0.2 1.5 0 21 12 2	0.8 1.3 2.4 3D 0.1 3.6 0.3 21 12 21	4 No 3 1	4.5 5.7 1D 7 0.1 2 21	0.5 0.8 3D 0.9 0.1 12 21	0.4 0.9 3D 1.5 0.1 12 21	0.4 0.6 0 3D 1.4 0.1 2 12 21	0.9 2.2 3D 2.1 0.1 12 21	2 2	1 2.3 3D 2.9 0.2 12 21	0.6 1.1 0 3D 1.3 0.1 1 12 21	0.8 1.2 (3D 1.4 0.1 1 12 21	0.7 1.6 3D 1.8 0.1 12 21	3D 1.4 0.1 12 21																	NUFVIU
PIJGIY	0.9 1.6 3D 1.1 0.1 14 21	0.9 1.5 3D 0.9 0.1 14 21	0.4 0.8 3D 2.1 0.2 12 21	0.6 1.1 3D 1.6 0.1 12 21	0.5 0.9 3D 0.1 1.9 0.1 12 21	0.5 1.1 1 3D 2.2 0.1 3 12 21 1 1	.5 3.5 1.2 3D 3E 3 0.1 0.7 14 21 14	2.1 0.7 1 3D 3D 1.4 0 21 14 2 2 14 2	1.3 1.8 3.1 3D 0.1 1.9 0.2 21 14 21	1 No 2 1	4.45.51D7.80.3221	0.4 0.8 3D 2.1 0.1 12 21	0.7 1.3 3D 1.6 0.1 14 21 21	0.6 1.2 0 3D 1.8 0.1 2 14 21	0.9 1.9 3D 2.8 0.1 14 21	No 	1.3 2.6 3D 1.3 0.1 14 21	0.5 1.2 0 3D 2.1 0.1 2 12 21	0.7 1.2 0 3D 2.7 0.2 1 12 21	0.9 1.6 3D 1.6 0 14 21	0.7 1.2 3D : 1.8 0.1 2.3 12 21 12	3D 0.2 21	٠															PIJGIY
PIJGOE	0.4 1 3D 1.3 0.1 14 21	0.3 0.8 3D 1.1 0.1 14 21	0.9 1.9 3D 2.2 0.2 12 21	0.6 1.5 3D 1.8 0.2 12 21	0.8 1.7 3D 2 0.2 12 21	0.9 2 1 3D 2.3 0.2 3 12 21 1	.3 2.7 0.4 3D 3E .2 0.2 1.1 14 21 14	0.6 0.5 1 3D 3D 1.6 0 21 14 2 14 2	1.3 0.9 1.6 3D 0.1 2.1 0.2 21 14 21	6 No 2 1	4.8 6.2 1D 7.9 0.4 2 21	0.9 1.9 3D 2.2 0.2 12 21	0.6 1.5 3D 1.8 0.1 14 21 21	0.7 1.7 1 3D 2 0.2 2 14 21	1.1 2.5 3D 2.8 0.2 14 21	(No 	0.5 1.1 3D 1.2 0 14 21	0.9 1.9 1 3D 2.2 0.2 2 12 21	.1 2.4 0 3D 2.8 0.2 1 12 21	0.6 1.5 3D 1.4 0.1 14 21	0.7 1.7 1 3D 2.1 0.1 2.4 12 21 12	2.1 3D 3 0.2 0.7 21 14	0.1 21															PIJGOE
POBRON	0.5 1.2 3D 0.7 0 14 21	0.4 1 3D 1.2 0.1 14 21	1 2 3D 1.7 0.2 12 21	0.8 1.6 3D 1.2 0.1 12 21	0.9 1.8 3D 1.4 0.1 12 21	1 2.1 1 3D 1 1.8 0.2 3 12 21 1	.4 2.8 0.5 3D 3E 5.4 0.2 1.2 14 21 14	1 0.6 1 3D 3D 3D 0.1 0.8 0 21 14 2	1.5 0.9 1.8 3D 0.1 2.1 0.2 21 14 21	3 No 2 1	4.8 6.2 1D 7.2 0.2 2 21	1 2 3D 1.6 0.2 12 21	0.7 1.6 3D 1.7 0.1 14 21	0.8 1.9 1 3D 1.2 0.1 14 21 ⁻	I.1 2.5 3D 3 0.1 14 21	(No 2 	0.5 1.1 3D 2.1 0.1 14 21	1 2 1 3D 1.9 0.1 1 12 21	.2 2.5 0 3D	0.5 1.3 3D 2.2 0.1 14 21	0.8 1.9 1.1 3D 3 1.4 0.1 2.2 12 21 12	2.1 0.3 3D 3 0.2 1.5 21 14	0.7 5 D 0.1 1. 21 1	3D .7 0.1 14 21														POBRON
POBRON01	0.3 0.7 3D 0.8 0.1 12 21 21	0.5 1.1 3D 1 0.1 12 21	0.8 1.4 3D 0.7 0.1 12 21	0.5 1 3D 0.4 0 12 21	0.7 1.2 3D 0.5 0 12 21 21	0.9 1.6 1 3D 0.9 0.1 3 12 21 1	.3 3.1 0.6 3D 3E .6 0.1 1.6 12 21 12	1.1 0.4 0 3D 3D 0.2 0.5 0.5 0.1 12 0.2 0.5	0.7 1 1.9 3D 0 2.8 0.3 21 12 21	9 No 3 1	4.4 5.7 1D 7.1 0.1 2 21	0.8 1.4 3D 0.6 0.1 12 21	0.7 1.5 3D 1.1 0.1 12 21	0.5 1.1 1 3D 0.6 0.1 2 12 21	1.2 2.7 3D 2.4 0.1 12 21	(No 2	0.8 1.9 3D 2.3 0.2 12 21	0.9 1.7 0 3D 1.1 0.1 1 12 21	0.9 1.7 0 3D 1.2 0.1 1 12 21	0.9 2 3D 1.6 0.1 12 21	0.6 1.2 1 3D 3 0.8 0.1 1.2 12 21 12	1.9 0.6 3D 3 0.1 1.6 21 12	1.4 0. SD 0.1 1. 21 1.	0.7 1.5 3D .8 0.2 1 12 21	3D 1.2 0.1 12 21												F	POBRON01
POVSIC	0.4 0.7 1D 0.2 2 21	0.5 0.9 1D 9.8 0.6 4 21	0.3 0.6 1D 9.4 0.5 4 21	0.1 0.4 1D 7.1 0.1 2 21	0.2 0.5 1D 9.5 0.5 4 21	0.4 0.8 1 1D 9.7 0.5 9 4 21 4	.4 3.2 0.8 1D 1E 0.8 0.5 7.8 4 21 2	1.4 0.2 0 1D 1D 0 0.3 7.2 0 21 2 2	0.5 1.4 2.4 1D 0.2 10.3 0.6 21 4 21	4 No 6	4.4 5.6 3D 1.9 0.1 14 21	0.3 0.5 1D 9.5 0.5 4 21	0.5 1 1D 9.7 0.6 4 21	0.2 0.6 1D 7.1 0.2 9 2 21	1 2.2 1D 9.4 0.6 4 21	(No - 1 1	0.9 2.1 1D 10.7 0.7 4 21	0.4 1 0 1D 9.7 0.5 6 4 21	0.6 1.1 (1D 0.7 0 1 2 21	0.7 1.4 1D 10.2 0.6 4 21	0.4 0.7 0.6 1D 7.3 0.1 9.2 2 21 4	1.1 0.6 1D 1 0.5 10.2 21 4	1.5 0. D 0.7 10 21 4	0.7 1.6 0 1D 0.3 0.7 7 4 21	0.5 1 1D 7.2 0.2 2 21	1D 7.2 0.1 2 21												POVSIC
SUJDAD	4.4 5.8 3D 1.4 0.1 14 21 21	4.9 8.5 3D 1.5 0.1 14 21	4.6 8.2 3D 2.5 0.2 12 21	4.3 5.7 3D 2 0.2 12 21 12 21	4.7 8.2 3D 2.3 0.2 12 21	4.8 8.4 4 3D 2.7 0.2 3 12 21 1	.8 8.4 4.6 3D 3E 5.1 0.2 0.8 14 21 14	6.2 4.3 5 0 3D 0.1 1.7 0 21 14 2	5.7 5.3 8.8 3D 0.1 1.7 0.2 21 14 21	3 No 2 1	0.8 1.7 1D 7.7 0.4 2 21	4.6 8.2 3D 2.5 0.2 12 21	4.8 8.4 3D 2.3 0.2 14 21 21	4.3 5.7 4 3D 2.2 0.2 3 14 21	3D 3.4 0.2 14 21	No 	5.3 9.3 3D 1.3 0.1 14 21	4.8 8.4 3D 2.7 0.2 12 21	4 5.4 5 3D 3 0.2 2 12 21	5.1 8.8 3D 2.3 0.1 14 21	4.4 5.8 4.4 3D 3 2.2 0.1 2.9 12 21 12	8 5.1 3D 3 0.3 1.1 21 14	8.8 5. 0.1 1. 21 1	5.1 8.8 4 3D .2 0.1 1 14 21 1	4.3 5.8 · 3D 1.4 0.1 14 21	4.3 5.7 3D 2 0.2 7. 12 21 2	3D B 0.4 21											SUJDAD
TUSHUL	0.6 1.2 3D 0 0.3 0 14 21	0.6 1.4 3D 0.8 0.1 14 21	1.2 2.2 3D 1.3 0.1 12 21	0.9 1.8 3D 0.8 0.1 12 21 21	1.1 2 3D 1.1 0.1 12 21 21	1.3 2.4 1 3D 1.5 0.1 3 12 21 1	.4 2.8 0.3 3D 3E 5.2 0.1 1.1 14 21 14	0.8 0.8 1 0 3D 0.1 0.6 21 14 2	1.5 0.7 1.8 3D 0 2.2 0.2 21 14 21	No 2 - - - - -	4.7 6.2 1D 7.2 0.2 2 21	1.2 2.2 3D 1.3 0.1 12 21 21	0.9 2.1 3D 1.3 0.1 14 21 21	1 1.9 1 3D 1 0.1 2 14 21 2	1.4 3.1 3D 2.6 0.1 14 21	No (0.5 1.2 3D 1.9 0.1 14 21	1.2 2.4 1 3D 1 1 1.6 0.1 1 12 21 1	.3 2.6 0 3D I.8 0.1 1 I2 21	0.9 2.1 3D 1.6 0.1 14 21	1 2 1.4 3D 3 1.1 0 1.8 12 21 12	2.5 0.5 3D 3 0.2 1.2 21 14	1 0. D 0.1 1. 21 1	0.5 1.1 (3D .4 0.1 (14 21	0.6 1.3 3D 0.7 0 14 21	0.9 1.8 4. 3D 0.8 0.1 7. 12 21 2	5 6.4 1D 3 0.2 1.4 21 14	3D 4 0.1 21										TUSHUL
UPULEX	0.1 0.3 1D 18.4 0.1 2 21	0.3 0.7 1D 18.9 0.2 2 21	0.6 1.1 1D 18 0 2 21	0.4 0.7 1D 18.1 0 2 21	0.5 0.9 1D 18.1 0 2 21	0.7 1.3 1 1D 18.3 0 1 2 21 2	.3 2.9 0.5 1D 1E 19 0.1 18.7 2 21 2	0.9 0.2 0 1D 10 10 0.2 18.2 21 2 2	0.5 1.1 2 1D 0 19.1 0.1 21 2 21	 No 1 1	4.5 5.7 No 	0.6 1.1 1D 18.1 0 2 21	0.5 1.1 1D 18.8 0.2 2 21	0.4 0.9 1D 18.3 0 11 2 21	1 2.3 1D 8.6 0 2 21	(No 1 	0.8 1.8 1D 19.5 0.2 2 21	0.7 1.4 0 1D 18.5 0 1 ¹ 2 21	0.8 1.6 (1D 7.6 0.1 1 2 21	0.7 1.5 1D 19.1 0.1 2 21	0.5 0.9 0.9 1D 18.4 0.1 17.9 2 21 2	1.5 0.5 1D 1 9 0 21 2	1.1 0. D 11 0.1 18 21 2	0.6 1.3 (D 1 8.9 0.1 2 2 21	0.3 0.7 1D 18 0 1 2 21	0.4 0.7 4. 1 D 8.2 0 - 2 21 -	4 5.8 0.0 No 1D - 18. - 2	5 1.2 1 3 0.1 18 21	D 3.4 0.1 2 21									UPULEX
VACZIJ	7.4 16.8 3D 1.3 0.1 12 21 21	7.5 17.3 3D 1.4 0.1 12 21	7.7 16.3 3D 0.6 0.1 12 21	7.5 16.5 3D 0.7 0 12 21 21	7.6 16.4 3D 0.6 0 12 21 21	7.816.693D0.90312211	16.3 7.3 3D 3E .8 0.1 2 .2 21 12	17.3 7.4 10 0 3D 0.2 1 0 21 12 2	6.6 9.8 16. 3D 0.1 3.2 0.3 21 12 21	5 No 3 1	1D 7 0.1 2 21	7.7 16.4 3D 0.6 0 12 21	7.7 17.2 3D 1.1 0.1 12 21	7.6 16.7 9 3D 0.8 0.1 2 12 21 ⁻	0.5 16 3D 2.3 0.1 12 21	No 2	9.9 16.8 3D 2.7 0.2 12 21	7.8 16.8 7 3D 1.1 0 1 12 21 7	7.3 16 9 3D 1.1 0.1 1 12 21	9.7 16.5 3D 1.8 0.1 12 21	7.6 16.8 7.8 3D 3 0.7 0.1 0.9 12 21 12	16.1 7.5 3D 3 0.1 2 21 12	17.4 7. 5 D 0.1 2. 21 1.	3D 2.2 0.2 12 21	7.2 16.5 3D 1.6 0.1 12 21	7.5 16.6 - 3D 0.7 0 9. 12 21 4	- 7.1 1D 4 0.5 2.4 21 12	1 16.9 7 3D 5 0.2 1 2 21 1	.4 16.8 3D .3 0.1 18 2 21	1D 8.1 0 2 21								VACZIJ
VOJLIQ	0.6 1.2 3D 0.6 0 14 21 21	0.6 1.2 3D 0.6 0 14 21	0.3 0.5 3D 1.2 0.1 12 21	0.3 0.7 3D 0.8 0.1 12 21	0.2 0.5 3D 1 0.1 12 21	0.4 0.8 1 3D 1.5 0.1 3 12 21 1	.4 3.5 1 3D 3E .2 0.1 1.1 14 21 14	1.8 0.4 0 3D 3D 0.1 0.6 0 21 14 2 2 14 2	0.9 1.5 2.8 3D 0.1 2.2 0.2 21 14 21	3 No 2 1	4.4 5.5 1D 7.3 0.1 2 21	0.3 0.5 3D 1.2 0.1 12 21	0.5 1 3D 1 0 14 21	0.3 0.7 0 3D 0.8 0.1 2 14 21	0.9 2.1 3D 2.3 0.1 14 21	No 2	1.1 2.4 3D 2.1 0.1 14 21	0.4 1 0 3D 1.4 0.1 1 12 21	0.6 0.9 0 3D 1.8 0.1 1 12 21	0.8 1.6 3D 1.6 0.1 14 21	0.3 0.7 0.4 3D 0.8 0 1.7 12 21 12	0.8 0.8 3D 3 0.2 1.3 21 14	1.9 0. 5 D 0.1 1. 21 1	0.9 2 0 3D .6 0.1 1 14 21	0.7 1.4 3D 1.1 0.1 14 21	0.3 0.6 4. 3D 0.8 0.1 7. 12 21 2	6 8.1 1.1 1D 3 0.2 1.1 21 14	1 2.2 0 3D 7 0.1 0 - 21 1	.6 1.1 7 3D .7 0 18 4 21	7.6 16.4 1D 8.5 0.1 2 21	3D 1.1 0.1 12 21							VOJLIQ
XEGTAF	0.3 0.5 No 	0.3 0.6 No 	0.7 1 No 	0.4 0.7 No 	0.6 0.9 No 	0.7 1.3 1 No 	.3 2.9 0.5 No No 	1 0.3 0 • No 	0.6 1 1.9 No 	1D 8.8 0.4 2 21	4.5 5.7 No 	0.7 1 No 	0.4 0.9 No 	0.4 0.7 C No 	0.9 2.1 No 	(1D 6.6 0.4 2 21	0.8 1.9 No 	0.7 1.2 0 No).9 1.5 (No 	0.6 1.4 No 	0.3 0.7 0.9 No I 	1.4 0.5 No N 	1.2 0. Io 	0.6 1.4 (No 	0.5 0.9 No 	0.4 0.6 4. No 	4 5.8 0. No 	7 1.5 0 No - -	.3 0.6 7 No 	7.5 16.9 (No 	0.5 1 No 	No 						XEGTAF
YIYSII	3D 1.7 0.1 14 21	3D 1.6 0.1 14 21	3D 2.4 0.2 12 21	3D 2.1 0.1 12 21	3D 2.2 0.1 12 21	3D 2.5 0.1 3 12 21 1	3D 3E 3.3 0.1 1.5 4 21 14	3D 0.1 2 0 21 14 2	3D 0.1 2.4 0.2 21 14 21	5.7 6.7 No 2 1	1D 7.9 0.4 2 21	3D 2.4 0.1 12 21	3D 2.1 0.1 14 21	3D 2.3 0.1 14 21	3D 30.1 1421	2.1 6.2 No 	3D 1.3 0.1 14 21	3D 2.4 0.1 3 12 21	3D 3.1 0.2 12 21	3D 2 0 14 21	3D 2.2 0.1 2.5 12 21 12	3D 3 0.2 1.3 21 14	 5 D 0 1. 21 1	3D .2 0 14 21	3D 2 0.1 14 21	3D 2 0.1 10 12 21 4	1D 3 0.7 1. ² 21 14	- 3D 7 0.1 1 - 21 1	3D .7 0.1 19 4 21	 1D 9.1 0.2 2 2 21	3D 2.4 0.1 12 21	3D 1.9 0.1 14 21	No 					YIYSII
YIYSOO	0.6 1.6 3D 2.3 0.1 14 21	0.5 1.5 3D 2 0.1 14 21	0.9 2.2 3D 2.8 0.2 12 21	0.7 1.9 3D 2.6 0.2 12 21	0.8 2.1 3D 2.7 0.2 12 21	0.9 2.3 1 3D 2.9 0.2 3 12 21 1	.3 3 0.6 3D 3E 3.7 0.2 1.7 14 21 14	1.3 0.7 1 3D 3D 1 0.1 2.4 0 21 14 2	1.8 1.2 2.1 3D 3D 3D 0.1 2.4 0.2 21 14 21	1 No 2 1	4.9 6.3 1D 8.6 0.3 2 21	0.9 2.2 3D 2.8 0.2 12 21	0.6 2 3D 2.5 0.1 14 21	0.8 2.1 1 3D 2.8 0.2 3 14 21 7	1.1 2.8 3D 3.3 0.2 14 21	(No 	0.6 1.2 3D 1.1 0.1 14 21	0.9 2.2 1 3D 2.8 0.2 3 12 21 7	.2 2.8 0 3D 3.5 0.3 1 12 21 5	0.7 1.9 3D 1.9 0.1 14 21	0.8 2.1 1 3D 2.7 0.2 2.9 12 21 12	2.3 0.4 3D 3 0.2 1.4 21 14	1.2 0. 5D 0.1 1. 21 1	0.5 1.1 0 3D .4 0 2 14 21	0.8 1.9 3D 2.6 0.2 14 21	0.7 1.9 5. 3D 2.5 0.2 11 12 21 4	1 8.9 0.8 1 D 1 0.6 2 21 14	3 1.5 0 3D 0.1 2 0.1 21 1	.6 1.6 9 3D .3 0.1 19 4 21	0.7 16.4 (1D 9.5 0.2 2 2 21	0.8 2.2 3D 2.8 0.2 12 21	0.6 1.7 3D 2.3 0.1 14 21	No 	3D 1.5 0.1 14 21				YIYSOO
ZIGZIY	0.9 2.1 3D 2 0.1 14 21	0.7 1.8 3D 2 0.1 14 21	1.1 2.6 3D 2.7 0.2 12 21	1 2.4 3D 2.4 0.2 12 21	1.1 2.5 3D 2.6 0.2 12 21	1.1 2.7 1 3D 3 2.8 0.2 3 12 21 1	.5 3.3 0.7 3D 3E 3.7 0.2 1.4 14 21 14	1.5 0.9 2 3D 3D 3D 0 2.2 0 21 14 2	2.3 1.1 2.1 3D 0.1 2 0.1 21 14 21	1 No 1	5.2 6.8 1D 8 0.3 2 21	1.1 2.6 3D 2.7 0.2 12 21	0.9 2.3 3D 2.5 0.1 14 21	1.1 2.6 1 3D 3 2.5 0.2 3 14 21 2	1.3 3.1 3D 3.6 0.2 14 21	0 No 	0.5 1 3D 1.7 0.1 14 21	1.1 2.5 1 3D 3 2.7 0.2 3 12 21 2	.4 3.2 (3D 3.1 0.2 2 12 21	0.6 1.8 3D 2.4 0.1 14 21	1 2.5 1.2 3D 3 2.5 0.1 3 12 21 12	2.7 0.5 3D 3 0.2 1.6 21 14	1.2 0. 5D 0.1 1. 21 1	0.5 1.3 3D .8 0.1 2 14 21	1 2.4 3D 2.2 0.1 14 21	0.9 2.3 5. 3D 2.3 0.2 10 12 21 4	5 9.6 0.8 1D .7 0.6 1.0 21 14	3 1.8 0 3D 3D 5 0.1 1 4 21 1	.9 2.1 9 3D 2 0.1 18 4 21	0.8 16.8 1D 8.9 0.1 2 2 21	1.1 2.6 3D 2.8 0.2 12 21	0.8 2.1 3D 2 0.1 14 21	 No 	0.6 1.4 3D 2.2 0.1 2 14 21 1	3D 2 0.1 14 21			ZIGZIY
β-CD·DMF	0.9 1.8 1D 8 0.4 4 21	0.8 1.8 1D 8.3 0.4 4 21	1.3 2.3 1D 7.9 0.6 4 21	1.1 2.1 1D 7.9 0.5 4 21	1.2 2.2 1D 7.9 0.5 4 21	1.4 2.5 1 1D 10 8 0.6 10 4 21 6	.7 3.2 0.6 2D 10 0.4 0.5 8.4 6 21 4	1.3 1 1 0 1D 1D 0.4 7.9 0 21 4 2	1.9 0.7 1.8 1D 0.4 8.6 0.3 21 4 21	B No 3 I	4.9 6.3 2D 8.9 0.4 8 21	1.3 2.3 1D 7.9 0.6 4 21	1.1 2.3 1D 8.3 0.5 4 21	1.1 2.2 1 1D 7.9 0.4 4 21	1.5 3.2 2D 10 0.4 6 21	(No - 8	0.6 1.6 1D 8.7 0.3 4 21	1.3 2.3 1 1D 1 1 8.1 0.6 7 4 21 1	.5 2.8 1D 7.5 0.5 8 4 21	1 2.2 1D 8.4 0.4 4 21	1.1 2.2 1.5 1D	2.5 0.7 1D 1 0.6 8.5 21 4	1.4 0. D 0.4 8. 21 4	0.7 1.7 0 1D 3.4 0.3 7 4 21	0.9 2 1D 7.8 0.3 4 21	1.1 2 5. 1D 7.9 0.5 9. 4 21 8	5 9.1 0.0 2D 4 0.4 8.3 21 4	5 1.5 0 1D 3 0.2 21 4	.9 1.8 7 1D 8 0.4 4 21	⁷ .4 17.4 No 	1.3 2.4 1D 7.8 0.5 4 21	0.8 1.8 1D 8.2 0.4 4 21	No	0.9 2 0. 1D 8.6 0.4 9 4 21 4	0.8 1.8 1D 9 0.4 8 4 21	1D 8.8 0.3 4 21		β-CD·DMF
	4 6.9 NIXN	4.2 7.1 VOX	4 6.8 000X3(4 6.8	4 6.8	4.1 6.8 5	NE 1.2		5.8 4.3 7.4 O O O O	4 XOOM	4.2 7.7 d0 22	4 6.8 00 22 X	4.2 7.1 YNNO	4 6.8 5 8 9 9 9	0.4 8.4	4	4.4 7.5 NOXL	4.1 7 3 NYBQ	8.7 6.4 4 X O Ø Ø	4.3 7.2 XHH	4.1 6.9 4 Ø U L	6.8 4.3	7.3 4. ≻⊔ טר	2 7.2 3 BOD	3.9 6.8 NO Ma	4 6.8 4.	4 8.3 4. Os	1 7.2 · OVD	4 6.9 TNHSI		4 6.7	4.1 7 OITC	GTAF	4.4 7.3 4.	00000000000000000000000000000000000000	4.3 7.7 ≻IZ9	D-DMF	
	AF	AF	BCL	BCI	BCI	BCI	BC	BU V	DE	DE	ā	GL	ЧH		X	ž	Я	LA	LA	Σ	ЯГ	z	ц.	Ē	РС	POE	Ĕ	ຮ	Ц	5	>	ž	XE	~	\geq	Ν	β-C	