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Dimethyl 2-(2,4,6-trimethoxybenzyl)-malonate

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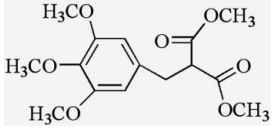
Key indicators: single-crystal X-ray study; T = 296 K; mean σ(C–C) = 0.002 Å; R factor = 0.040; wR factor = 0.112; data-to-parameter ratio = 18.8.

In the title compound, $C_{15}H_{20}O_7$, the benzene ring makes dihedral angles of 69.17 (5) and 80.81 (4)° with the two side chains of malonate. The two malonate side chains comprising C/C/O/C atoms are oriented at right angles [86.26 (6)°] with respect to each other. In the crystal structure, the crystal packing is stabilized by weak non-classical intermolecular C—H···O hydrogen bonds, which link the molecules into an infinite network.

Related literature

Substituted malonate, an important organic intermediate, is electrooxidized in methanol in the presence of halogen ions to afford the corresponding halomalonates, see: Okimoto & Takahashi (2002). For a related structure, see: Liu *et al.* (2010).



Experimental

Crystal data

$C_{15}H_{20}O_7$ M_r = 312.31
Monoclinic, $P2_1/n$
a = 11.6173 (15) Å
b = 8.1192 (10) Å
c = 17.236 (2) Å
β = 103.968 (2)°
V = 1577.7 (3) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.11 mm^{-1}
T = 296 K
0.28 × 0.24 × 0.20 mm

Data collection

Rigaku Saturn CCD area-detector diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2005)
T_{min} = 0.904, T_{max} = 0.935
9686 measured reflections
3851 independent reflections
2969 reflections with $I > 2\sigma(I)$
R_{int} = 0.019

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.040
$wR(F^2)$ = 0.112
S = 1.05
3851 reflections
205 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.21 e Å^{-3}
$\Delta\rho_{min}$ = −0.15 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
C5—H5b···O2^{i}	0.96	2.54	3.353 (2)	142
C5—H5a···O5^{ii}	0.96	2.57	3.503 (2)	164
C3—H3···O6^{iii}	0.98	2.54	3.499 (2)	168

Symmetry codes: (i) $-x-\frac{1}{2}, y-\frac{1}{2}, -z+\frac{1}{2}$; (ii) $-x+\frac{1}{2}, y-\frac{1}{2}, -z+\frac{1}{2}$; (iii) $-x+\frac{1}{2}, y+\frac{1}{2}, -z+\frac{1}{2}$.

Data collection: *CrystalClear* (Rigaku/MSC, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

This work was supported by the NSFC (grant No.30873139).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2277).

References

Liu, S.-X., Zhang, Z.-H., Yang, Y.-F., Zhen, X.-L. & Han, J.-R. (2010). *Acta Cryst.* E**66**, o1383.
Okimoto, M. & Takahashi, Y. (2002). *Synthesis*, **15**, 2215–2219.
Rigaku/MSC (2005). *CrystalClear*. Rigaku/MSC Inc., The Woodlands, Texas, USA.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supplementary materials

Acta Cryst. (2010). E**66**, o1384 [**doi:10.1107/S1600536810015928**]

Dimethyl 2-(2,4,6-trimethoxybenzyl)malonate

S.-X. Liu, X. Lu, S.-R. Gao, J.-R. Han and X.-L. Zhen

Comment

Substituted malnoate is a very important organic intermediate. It is electrooxidized in methanol in the presence of halogen ions to afford the corresponding halomalonates (Okimoto & Takahashi, 2002). We have synthesized dimethyl 2-(quinolin-methyene)malonate, and reported its structure (Liu *et al.*, 2010). We now report the synthesis and structure of the title compound, (I).

In the title compound (Fig. 1), the benzene ring makes dihedral angles of 69.17 (5) and 80.81 (4)° with two side chains of malonate. In the crystal structure, the two malonate side chains comprising C/C/O/O atoms (C2/C3/O1/O2 and C113/C4/O3/C4) are oriented at right angles (86.26 (6)°) with respect to each other. In the crystal structure, the crystal packing is stabilized by weak non-classical intermolecular C—H···O hydrogen bonds which link the molecules into an infinite network; details have been provided in Table 1 and Fig. 2.

Experimental

An anhydrous methanol solution (130 ml) of 1-brommethyl-3,4,5-trimethoxy (26.0 g, 0.1 mol) was added to an anhydrous methanol solution (180 ml) of sodium methoxide (5.4 g, 0.1 mol) and dimethyl malonate (26.4 g, 0.2 mol). The mixture was refluxed for 6 hours. The product was isolated with silica gel column, then the solvent was removed and added petroleum ether (5 ml) to the white precipitate. The precipitate was then isolated, recrystallized from n-hexane, and dried in a vacuum to give the title compound in 75% yield. Colorless single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an n-hexane solution.

Refinement

The H atoms were included in calculated positions (C—H = 0.93–0.98 Å) and refined as riding with U_{iso}(H) = 1.2U_{eq}(C) or 1.5U_{eq}(methyl C).

Figures

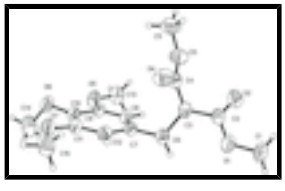


Fig. 1. The title structure plotted with displacement ellipsoids at 50% probability level.

Fig. 2. A unit cell showing intermolecular hydrogen bonds by dashed lines; the H-atoms not involved in H-bonds have been excluded for clarity.

Dimethyl 2-(2,4,6-trimethoxybenzyl)malonate

Crystal data

$C_{15}H_{20}O_7$	$F(000) = 664$
$M_r = 312.31$	$D_x = 1.315$ Mg m^{-3}
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -P 2yn	Cell parameters from 3043 reflections
$a = 11.6173$ (15) Å	$\theta = 2.4$–$24.5°$
$b = 8.1192$ (10) Å	$\mu = 0.11$ mm^{-1}
$c = 17.236$ (2) Å	$T = 296$ K
$\beta = 103.968$ (2)°	Prism, colourless
$V = 1577.7$ (3) Å^3	0.28 × 0.24 × 0.20 mm
$Z = 4$	

Data collection

Rigaku Saturn CCD area-detector diffractometer	3851 independent reflections
Radiation source: fine-focus sealed tube	2969 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.019$
Detector resolution: 7.31 pixels mm^{-1}	$\theta_{max} = 28.3°$, $\theta_{min} = 1.9°$
ω and φ scans	$h = -14\rightarrow12$
Absorption correction: multi-scan (CrystalClear; Rigaku/MSC, 2005)	$k = -8\rightarrow10$
$T_{min} = 0.904$, $T_{max} = 0.935$	$l = -19\rightarrow22$
9686 measured reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.040$	H-atom parameters constrained
$wR(F^2) = 0.112$	$w = 1/[\sigma^2(F_o^2) + (0.0481P)^2 + 0.3127P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	$(\Delta/\sigma)_{max} < 0.001$
3851 reflections	$\Delta\rho_{max} = 0.21$ e Å^{-3}
205 parameters	$\Delta\rho_{min} = -0.15$ e Å^{-3}
0 restraints	Extinction correction: SHELXL97 (Sheldrick, 2008), $Fc^* = kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Primary atom site location: structure-invariant direct methods

Extinction coefficient: 0.0172 (18)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor wR and goodness of fit *S* are based on F^2, conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) etc. and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
C1	−0.13204 (15)	0.9367 (2)	0.43372 (11)	0.0635 (4)
H1A	−0.2035	0.9372	0.3918	0.095*
H1B	−0.1042	1.0476	0.4447	0.095*
H1C	−0.1479	0.8893	0.4811	0.095*
C2	−0.07201 (11)	0.68523 (16)	0.39019 (7)	0.0404 (3)
C3	0.02668 (11)	0.59332 (15)	0.36509 (7)	0.0395 (3)
H3	0.0329	0.6362	0.3131	0.047*
C4	−0.01013 (11)	0.41471 (17)	0.35462 (8)	0.0421 (3)
C5	−0.10242 (17)	0.2143 (2)	0.26180 (11)	0.0722 (5)
H5A	−0.0371	0.1390	0.2762	0.108*
H5B	−0.1390	0.2036	0.2058	0.108*
H5C	−0.1595	0.1898	0.2922	0.108*
C6	0.14750 (11)	0.61262 (17)	0.42412 (8)	0.0424 (3)
H6A	0.1692	0.7282	0.4279	0.051*
H6B	0.1413	0.5765	0.4766	0.051*
C7	0.24446 (11)	0.51492 (15)	0.39988 (8)	0.0389 (3)
C8	0.26120 (11)	0.52716 (16)	0.32288 (8)	0.0403 (3)
H8	0.2142	0.5980	0.2861	0.048*
C9	0.34815 (11)	0.43345 (15)	0.30108 (8)	0.0390 (3)
C10	0.41922 (11)	0.32810 (16)	0.35617 (8)	0.0411 (3)
C11	0.40304 (11)	0.31759 (16)	0.43313 (8)	0.0429 (3)
C12	0.31541 (11)	0.41072 (16)	0.45482 (8)	0.0426 (3)
H12	0.3044	0.4030	0.5064	0.051*
C13	0.30064 (15)	0.5376 (2)	0.16826 (9)	0.0598 (4)
H13A	0.2188	0.5060	0.1589	0.090*
H13B	0.3260	0.5284	0.1194	0.090*
H13C	0.3097	0.6494	0.1868	0.090*
C14	0.61684 (16)	0.2734 (3)	0.35421 (15)	0.0899 (7)
H14A	0.6255	0.3796	0.3318	0.135*
H14B	0.6648	0.1946	0.3348	0.135*
H14C	0.6419	0.2792	0.4114	0.135*

C15	0.47068 (17)	0.2007 (2)	0.56275 (10)	0.0675 (5)
H15A	0.4958	0.3047	0.5876	0.101*
H15B	0.5227	0.1153	0.5893	0.101*
H15C	0.3911	0.1778	0.5665	0.101*
O1	−0.04280 (9)	0.84060 (12)	0.40905 (6)	0.0532 (3)
O2	−0.16628 (8)	0.62656 (13)	0.39094 (6)	0.0558 (3)
O3	−0.06001 (10)	0.38021 (14)	0.27852 (6)	0.0579 (3)
O4	0.00316 (11)	0.31615 (13)	0.40747 (6)	0.0613 (3)
O5	0.37059 (9)	0.43260 (12)	0.22680 (6)	0.0505 (3)
O6	0.49714 (9)	0.22459 (13)	0.33151 (6)	0.0539 (3)
O7	0.47421 (10)	0.20690 (14)	0.48168 (6)	0.0623 (3)

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0574 (9)	0.0518 (9)	0.0869 (12)	0.0155 (7)	0.0283 (8)	−0.0038 (8)
C2	0.0369 (6)	0.0443 (7)	0.0404 (6)	0.0060 (5)	0.0103 (5)	0.0068 (5)
C3	0.0389 (6)	0.0423 (7)	0.0400 (6)	0.0047 (5)	0.0150 (5)	0.0049 (5)
C4	0.0364 (6)	0.0462 (7)	0.0450 (7)	0.0040 (5)	0.0122 (5)	0.0005 (6)
C5	0.0692 (11)	0.0745 (12)	0.0693 (11)	−0.0141 (9)	0.0099 (9)	−0.0216 (9)
C6	0.0380 (6)	0.0429 (7)	0.0484 (7)	0.0028 (5)	0.0147 (5)	−0.0035 (6)
C7	0.0342 (6)	0.0365 (6)	0.0473 (7)	−0.0008 (5)	0.0125 (5)	−0.0037 (5)
C8	0.0371 (6)	0.0375 (6)	0.0481 (7)	0.0036 (5)	0.0136 (5)	0.0034 (5)
C9	0.0380 (6)	0.0365 (6)	0.0451 (7)	−0.0021 (5)	0.0150 (5)	−0.0022 (5)
C10	0.0377 (6)	0.0360 (6)	0.0505 (7)	0.0036 (5)	0.0124 (5)	−0.0061 (5)
C11	0.0420 (7)	0.0383 (7)	0.0467 (7)	0.0038 (5)	0.0078 (5)	−0.0010 (5)
C12	0.0432 (7)	0.0438 (7)	0.0422 (7)	0.0014 (5)	0.0129 (5)	−0.0018 (5)
C13	0.0678 (10)	0.0642 (10)	0.0507 (8)	0.0134 (8)	0.0206 (7)	0.0094 (7)
C14	0.0486 (10)	0.1037 (16)	0.1212 (18)	0.0205 (10)	0.0282 (10)	−0.0129 (13)
C15	0.0708 (11)	0.0756 (12)	0.0525 (9)	0.0189 (9)	0.0081 (8)	0.0119 (8)
O1	0.0482 (5)	0.0417 (5)	0.0760 (7)	0.0066 (4)	0.0271 (5)	0.0019 (5)
O2	0.0393 (5)	0.0585 (6)	0.0730 (7)	0.0012 (4)	0.0202 (5)	−0.0026 (5)
O3	0.0628 (6)	0.0628 (7)	0.0462 (6)	−0.0067 (5)	0.0098 (5)	−0.0055 (5)
O4	0.0764 (8)	0.0473 (6)	0.0553 (6)	−0.0060 (5)	0.0061 (5)	0.0080 (5)
O5	0.0555 (6)	0.0529 (6)	0.0487 (5)	0.0127 (4)	0.0238 (4)	0.0057 (4)
O6	0.0521 (6)	0.0520 (6)	0.0598 (6)	0.0174 (4)	0.0176 (5)	−0.0061 (5)
O7	0.0716 (7)	0.0625 (7)	0.0521 (6)	0.0302 (6)	0.0134 (5)	0.0090 (5)

Geometric parameters (Å, °)

C1—O1	1.4411 (17)	C8—C9	1.3869 (17)
C1—H1A	0.9600	C8—H8	0.9300
C1—H1B	0.9600	C9—O5	1.3668 (15)
C1—H1C	0.9600	C9—C10	1.3915 (18)
C2—O2	1.1973 (16)	C10—O6	1.3756 (15)
C2—O1	1.3262 (17)	C10—C11	1.3864 (19)
C2—C3	1.5159 (17)	C11—O7	1.3630 (16)
C3—C4	1.5104 (19)	C11—C12	1.3898 (18)
C3—C6	1.5298 (18)	C12—H12	0.9300

C3—H3	0.9800	C13—O5	1.4175 (18)
C4—O4	1.1941 (16)	C13—H13A	0.9600
C4—O3	1.3296 (16)	C13—H13B	0.9600
C5—O3	1.439 (2)	C13—H13C	0.9600
C5—H5A	0.9600	C14—O6	1.408 (2)
C5—H5B	0.9600	C14—H14A	0.9600
C5—H5C	0.9600	C14—H14B	0.9600
C6—C7	1.5165 (17)	C14—H14C	0.9600
C6—H6A	0.9700	C15—O7	1.4086 (19)
C6—H6B	0.9700	C15—H15A	0.9600
C7—C12	1.3839 (18)	C15—H15B	0.9600
C7—C8	1.3907 (18)	C15—H15C	0.9600
O1—C1—H1A	109.5	O5—C9—C8	124.92 (12)
O1—C1—H1B	109.5	O5—C9—C10	114.85 (11)
H1A—C1—H1B	109.5	C8—C9—C10	120.22 (12)
O1—C1—H1C	109.5	O6—C10—C11	120.62 (12)
H1A—C1—H1C	109.5	O6—C10—C9	119.40 (12)
H1B—C1—H1C	109.5	C11—C10—C9	119.70 (11)
O2—C2—O1	123.81 (12)	O7—C11—C10	115.20 (11)
O2—C2—C3	124.33 (12)	O7—C11—C12	124.66 (12)
O1—C2—C3	111.83 (11)	C10—C11—C12	120.08 (12)
C4—C3—C2	107.15 (10)	C7—C12—C11	120.16 (12)
C4—C3—C6	111.45 (10)	C7—C12—H12	119.9
C2—C3—C6	113.29 (10)	C11—C12—H12	119.9
C4—C3—H3	108.3	O5—C13—H13A	109.5
C2—C3—H3	108.3	O5—C13—H13B	109.5
C6—C3—H3	108.3	H13A—C13—H13B	109.5
O4—C4—O3	123.82 (13)	O5—C13—H13C	109.5
O4—C4—C3	124.86 (12)	H13A—C13—H13C	109.5
O3—C4—C3	111.32 (11)	H13B—C13—H13C	109.5
O3—C5—H5A	109.5	O6—C14—H14A	109.5
O3—C5—H5B	109.5	O6—C14—H14B	109.5
H5A—C5—H5B	109.5	H14A—C14—H14B	109.5
O3—C5—H5C	109.5	O6—C14—H14C	109.5
H5A—C5—H5C	109.5	H14A—C14—H14C	109.5
H5B—C5—H5C	109.5	H14B—C14—H14C	109.5
C7—C6—C3	112.73 (10)	O7—C15—H15A	109.5
C7—C6—H6A	109.0	O7—C15—H15B	109.5
C3—C6—H6A	109.0	H15A—C15—H15B	109.5
C7—C6—H6B	109.0	O7—C15—H15C	109.5
C3—C6—H6B	109.0	H15A—C15—H15C	109.5
H6A—C6—H6B	107.8	H15B—C15—H15C	109.5
C12—C7—C8	119.96 (11)	C2—O1—C1	115.31 (11)
C12—C7—C6	119.38 (11)	C4—O3—C5	116.19 (13)
C8—C7—C6	120.66 (11)	C9—O5—C13	117.23 (11)
C9—C8—C7	119.87 (12)	C10—O6—C14	114.95 (12)
C9—C8—H8	120.1	C11—O7—C15	118.30 (12)
C7—C8—H8	120.1		

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C5—H5b···O2^{i}	0.96	2.54	3.353 (2)	142
C5—H5a···O5^{ii}	0.96	2.57	3.503 (2)	164
C3—H3···O6^{iii}	0.98	2.54	3.499 (2)	168

Symmetry codes: (i) −*x*−1/2, *y*−1/2, −*z*+1/2; (ii) −*x*+1/2, *y*−1/2, −*z*+1/2; (iii) −*x*+1/2, *y*+1/2, −*z*+1/2.

Fig. 1

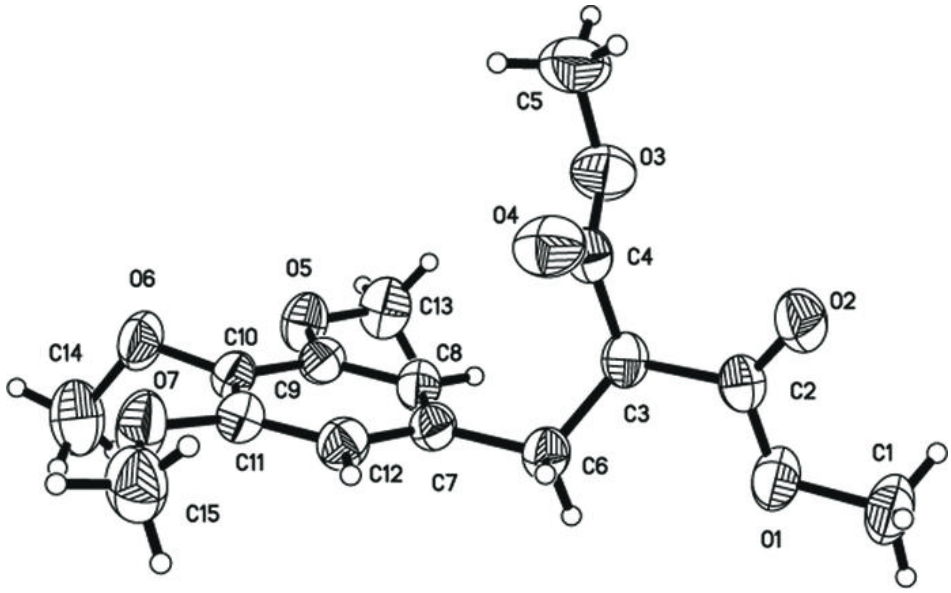
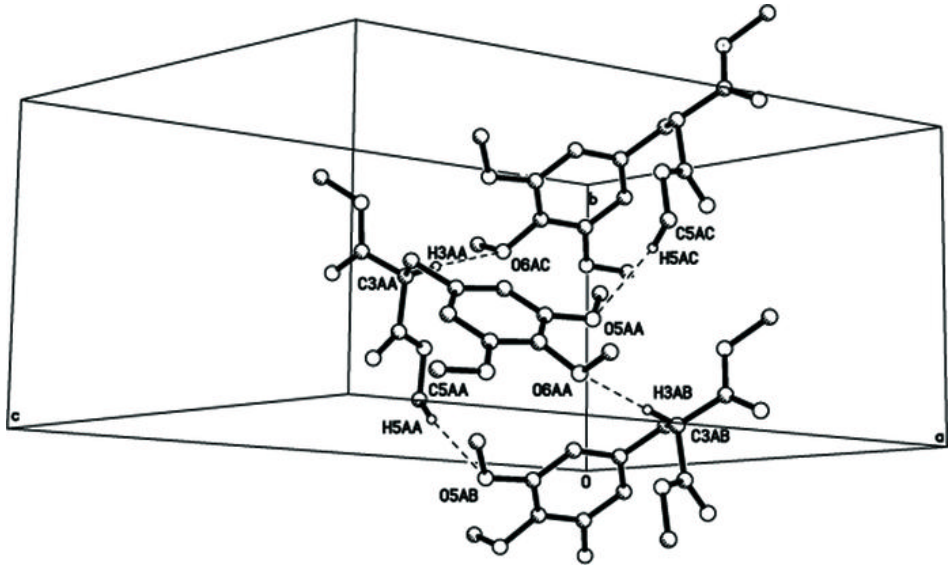

C5
O3
O4
C4
O2
C13
O5
O6
C10
C14
C8
O7
C9
C3
C2
C1
C11
C12
C7
C6
C15
O1

Fig. 2


H3AA
O6AC
C5AC
H5AC
C3AA
O5AA
C5AA
O6AA
H3AB
H5AA
C3AB
O5AB
c
a
b
0