## AN APPROACH TO BIS(AMINO ACID)S UTILIZING DIMETHYL

#### 2,4-BIS(DIAZO)-3-OXOGLUTARATE AND STUDIES OF

#### TRIS(2,6-DIHYDROXYPHENYL)E, E = B, P.

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#### THESIS ABSTRACT

# AN APPROACH TO BIS(AMINO ACID)S UTILIZING DIMETHYL 2,4-BIS(DIAZO)-3-OXOGLUTARATE AND STUDIES OF TRIS(2,6-DIHYDROXYPHENYL)E, E = B, P.

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When treated with a diazo transfer agent, dimethyl 3-oxoglutarate afforded a high yield of the bis(diazo) compound dimethyl 2,4-bis(diazo)-3-oxoglutarate, 33. The bis(diazo) compound, catalyzed by Rh<sub>2</sub>(OAc)<sub>4</sub>, reacted with an excess of O-benzyl carbamate, (CbzNH<sub>2</sub>), to give the product of double N-H insertion, 36, [MeO<sub>2</sub>C-CH(NHCbz)]<sub>2</sub>C=O. Thus in two steps one obtains a protected bis(amino acid). Compound 36 exhibits complex <sup>1</sup>H and <sup>13</sup>C NMR spectra, which may be rationalized by assuming 36 is predominantly enolic.

Attempted removal of the Cbz protecting groups gave startling results. Hydrogenation of 36 over Pd/C led to dimethyl 2-carboxamidosuccinate, 44,  $MeO_2CCH_2CH(CONH_2)(CO_2Me)$ , the structure of which was proven by x-ray crystallography.

In other studies, tris(2,6-dimethoxyphenyl)boron was synthesized by literature procedures. Demethylation was attempted using AlCl<sub>3</sub>/toluene, and BBr<sub>3</sub>·SMe<sub>2</sub>. In both cases, resorcinol, the product of B-C bond cleavage was observed.

Tris(2,6-dihydroxyphenyl)phosphonium chloride 63 was synthesized and its x-ray crystal structure was determined. It binds ethyl ether via a hydrogen bond. The three aryl rings adopt propeller symmetry, with torsion angles relative to the P-H direction of  $20.5^{\circ}$ ,  $24.5^{\circ}$ , and  $30.8^{\circ}$ . Compared to the average C-P-H angle of  $[Ph_3PH]^+$  -  $108.2^{\circ}$ , the average C-P-H angle of 63 is smaller, namely  $104.9^{\circ}$ .

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#### 1. PART I. INTRODUCTION

#### **1.1 Introduction to Diazocarbonyl Compounds**

The date of the first recorded synthesis of an  $\alpha$ -diazocarbonyl compound is from the work of Curtius<sup>1,2</sup> on diazotization of natural  $\alpha$ -amino acids (ethyl diazoacetate was first synthesized in 1883 from glycine). The simple diazocarbonyl compounds became available only in the late 1920s with the work of Arndt and Eistert,<sup>3-5</sup> and of Bradley and Robinson.<sup>6</sup> Now, modern organic synthesis continues to benefit from the unique versatility of diazocarbonyl compounds in cyclopropanation, X-H insertion (X = C, N, O, S, Se, P, halogen), Wolff rearrangement, ylide formation with subsequent transformations, aromatic cycloaddition and substitution, and many other useful reactions.

There are perhaps four principal reasons for the high level of activity in the diazo compound area. First, the enormous number of transformations that can occur with diazocarbonyl compounds makes them extremely versatile reactants. Second, methodology for their synthesis has continued to develop on a broad front, and there are now available well-tested, reliable procedures for the preparation of all the main classes of diazocarbonyl compounds. Third, the introduction in the 1970s of dirhodium(II) catalysts for diazocarbonyl decomposition opened up numerous new opportunities for highly chemoselective transformations that were largely inaccessible with conventional copper catalysts. Fourth, the ever-increasing demands for stereocontrol in the production of molecules of high enantiopurity have led to the introduction of chiral catalysts for asymmetric transformations of diazocarbonyl compounds.<sup>7</sup>

The synthesis of diazoketones involves addition of an acyl chloride to ethereal diazomethane<sup>3-5</sup> at or below 0  $^{\circ}$ C (eq. 1). Numerous synthetic intermediates containing the diazoketone functional group have been obtained in this way.

$$RCOCI + 2 CH_2N_2 \xrightarrow{Et_2O} RCOCHN_2 + CH_3CI + N_2$$
(1)

#### **1.2 Diazo Transfer Reaction**

Diazo transfer is now the standard route which is widely used to transfer a complete diazo group from a donor to an acceptor. Since the middle of the last century, sulfonyl azide compounds have been developed to serve as excellent diazo donors.<sup>8</sup> The diazo group transfers to the  $\alpha$ -methylene position of a carbonyl compound by the simple diazo transfer procedure of direct exposure to tosyl azide in dry acetonitrile or ethanol using triethylamine as a base (eq. 2).

The mechanism of diazo transfer reaction has been developed by M. Regitz Saarbrücken (Scheme 1).<sup>9</sup>



Scheme 1

Although tosyl azide has been by far the most frequently employed diazo donor, in their synthesis of and was used by Doering and DePuy in 1953 diazocyclopentadiene,<sup>10</sup> work in the Merck laboratories<sup>11</sup> has raised doubts regarding the safety aspects of this reagent. With these worries, Merck chemists<sup>11,12</sup> have examined sulforyl azides 1 to 12 from the standpoint of utility, thermal stability, ease of handling, and safety. It was found that methanesulfonyl (mesyl) azide 1 was the most hazardous and dangerous diazo transfer reagent of the group. Compared to more dangerous transfer reagent, tosyl azide, p-acetamidobenzenesulfonyl azide (p-ABSA) 7 offers more safety, yield, and ease of manipulation in diazo transfer. This reagent is currently commercially available and has some advantage in safety and yield, and so it is used in our laboratory for the diazo transfer reactions. These diazo transfer reagents, 1 to 12, are very efficient, giving good yield (>80%) in various diazo transfer reactions.



These diazo transfer reagents may be made from the corresponding sulfonyl chlorides (eq. 3).

/n

11

12

C<sub>9</sub>H<sub>19</sub>

10

Br

9

$$\begin{array}{c} SO_2CI \\ \downarrow \\ HN \\ O \end{array} + NaN_3 \xrightarrow{TBAB^a} CH_2CI_2 \\ HN \\ O \end{array} \begin{array}{c} SO_2N_3 \\ \downarrow \\ HN \\ CH_3 \end{array}$$
(3)

<sup>a</sup> tetrabutylammonium bromide

With a diazo transfer reagent such as 7, the  $\alpha$ -diazocarbonyl compound is readily formed. When the reaction site is activated by *two* flanking carbonyls, the diazo transfer reaction works even more successfully. For example, Hazen et al.<sup>11</sup> employed the naphthalene-based diazo transfer reagent **4** in the high-yield synthesis of dimethyl diazomalonate, DDM (eq. 4).

$$H_{3}CO \longrightarrow OCH_{3} + 4 \xrightarrow{Et_{3}N} H_{3}CO \longrightarrow H_{3}CO \longrightarrow OCH_{3} (4)$$

$$DDM, 95\%$$

#### **1.3 Catalysts for Metal Carbene Transformations**

Diazo compounds are inherently unstable to acid-promoted decomposition. Transition metals, which act as Lewis acids, are therefore effective catalysts for diazo decomposition.<sup>14</sup> The reactivities of diazocarbonyl compounds and alkyl/aryl diazo compounds are different. This may be rationalized on the basis of the pK<sub>a</sub> values of the corresponding diazonium ions. As an example, diazomethane is the conjugate base of the methanediazonium ion (eq. 5), whose p $K_a$  value<sup>14</sup> of 10 suggests the driving force for the

high reactivity of diazomethane towards Brønsted acids, and, by extension, Lewis acids.

$$\begin{array}{c} H \\ H - C - N \equiv N \\ H \end{array} \xrightarrow{} \begin{array}{c} H \\ C = N = N \end{array} \xrightarrow{} \begin{array}{c} H \\ C = N = N \end{array} \xrightarrow{} \begin{array}{c} H \\ C = N = N \end{array} \xrightarrow{} \begin{array}{c} H \\ H \end{array} \xrightarrow{} \begin{array}{c} H \\ \end{array} \xrightarrow{} \begin{array}{c} H \\ H \end{array} \xrightarrow{} \begin{array}{c} H \\ \end{array}$$

In contrast, the p $K_a$  values of diazonium ions derived from diazoesters (**a**) and diazoketones (**b**) have been estimated to be between -5 and -2, respectively.<sup>15</sup> This is



consistent with the much greater stability of diazoesters and diazoketones toward acid-promoted decomposition. So, various transition metals have been developed as diazo decomposition catalysts to afford more efficient reactivity in diazo decomposition reactions.

# 1.3.1 Mechanism of catalytic diazo decomposition. Metal carbene generation and reactions

Transition metal complexes that are effective catalysts for diazo decomposition are Lewis acids.<sup>13</sup> Their catalytic activity depends on coordinative unsaturation at the

metal center, which allows them to react as electrophiles with diazo compounds.<sup>7</sup> In the generally accepted mechanism for catalytic decomposition of diazo compounds (Scheme 2),<sup>13,16-19</sup> electrophilic addition causes the loss of dinitrogen and production of a metal-stabilized carbene (**14**). Transfer of the electrophilic carbene entity to an electron-rich substrate (S:) regenerates the catalytically active  $L_nM$  and completes the catalytic cycle.



Scheme 2

The activities of catalytically active transition metal compounds towards diazo decomposition are dependent on both the electrophilicity of the transition metal compound and on the stability of the diazo compound.<sup>15</sup> Among diazocarbonyl compounds, those with two carbonyl groups flanking the diazo-bearing carbon are more stable toward transition metal-catalyzed decomposition than those with only one

carbonyl group.<sup>17</sup> Diazoesters are generally more stable than diazoketones, and diazoamides are more stable than diazoesters. This profile is a useful guide to determining the reaction conditions required to generate a metal carbene. For example, diazoacetoacetates and diazomalonates require higher temperatures for reactions with transition metal catalysts than do diazoacetates, which can undergo catalytic nitrogen loss at or below room temperature.<sup>15</sup>



#### **1.3.2 Rhodium catalysts for diazo decomposition**

Dirhodium(II) catalysts are the most effective and versatile for diazo decomposition.<sup>7,16-20</sup> Dirhodium(II) tetraacetate (**15**) was first introduced by Teyssié and



co-workers in 1973. Control of reactivity and selectivity in such catalysts can be provided by varying the catalysts' bridging carboxylate (e.g.  $16^{21}$ ) or carboxamide (e.g.  $17^{22}$ ) ligands (in the structures above, the ligand shown is repeated three more times). Rhodium acetate 15 was originally employed as a catalyst for diazo decomposition in the presence of aniline (eq. 6), namely, insertion of a metal-carbene into an N-H bond. The reaction afforded the desired product 18 in 70% yield.<sup>23</sup> Compared to copper catalyst with much lower yield (<30% usually),<sup>24</sup> the dirhodium catalyst offers the advantages of mild conditions and higher yield.



#### 1.4 N-H insertion

Metal carbenes derived from  $\alpha$ -diazocarbonyl compounds, e.g. **19**, are highly electrophilic. They readily react with an available Lewis base (B:) to form a zwitterionic adduct (Scheme 3, in which B: is PhNH<sub>2</sub>). Catalytically generated electrophilic metal



carbenes add to Lewis bases, characteristically heteroatom-substituted organic compounds, to form zwitterionic adducts **20**, which can either dissociate to form rearranged products (or form a "free" ylide) and the catalyst, or revert to the metal carbene and Lewis base. With catalytically generated metal carbenes, especially those of rhodium, the metal-carbon bond of the intermediate metal-stabilized ylide is generally weaker than the R<sub>2</sub>C-base bond, and the preferred cleavage is that of the metal-carbon bond. The ease with which catalytically generated metal carbenes transfer the carbene entity to a heteroatom of an organic base is the basis for the synthetic utility to this diazo decomposition methodology.<sup>7</sup>



Scheme 3

#### **1.4.1 Intermolecular N-H Reaction**

Insertion into N-H bonds by diazocarbonyl compounds attracted little attention as a synthetic route to  $\alpha$ -amino ketones or esters until 1978, when a bicyclic  $\beta$ -lactam synthesis was published from the Merck laboratories.<sup>25-27</sup> Since then the power of this reaction has been demonstrated dramatically, especially the intramolecular reaction leading to nitrogen heterocycles. As mentioned, the introduction of rhodium(II) catalysts for N-H insertion led to major improvements with reaction between EDA and aniline in the presence of rhodium(II) acetate furnishing in 70% yield the insertion product (eq. 6).<sup>28</sup>

Reactions are carried out either neat or in solution (benzene or ethylene glycol dimethyl ether). In a similar fashion, Landais and Planchenault<sup>29</sup> used rhodium(II) acetate in benzene to bring about N-H insertion of aniline in an allylic amino ester

synthesis (eq. 7).



N-H insertion with diazocarbonyl compounds is not limited to amino functions. Insertion reactions involving amides,  $\beta$ -lactams, and carbamates are also known. Some examples are shown in eqs. 8,<sup>30</sup> 9,<sup>31</sup> and 10.<sup>30</sup> For the reaction in eq. 10 two different



N-H sites are available, one in the form of an amide and the other a carbamate. The former is preferred.

#### **1.4.2 Intramolecular N-H Reaction**

By far the most successful metal-catalyzed N-H insertion reactions have been intramolecular, leading to nitrogen heterocycles. Most recent studies, mainly by Rapoport and co-workers,<sup>32</sup> have established that while five-membered formation is preferred kinetically in the intramolecular insertion, it is possible to construct four- to six-membered aza rings from simple, conformationally mobile acyclic diazoester precursors. Rhodium(II) acetate was used as catalyst throughout, and variations in solvent, temperature, and catalyst concentration were found to play a role in determining product distributions. A summary of the results is shown in Scheme 4. Cyclization to four- and five-membered rings containing N occurred efficiently and selectively (entries 1 and 2). With the diazo compound in entry 3, the opportunity exists for both intramolecular C-H and N-H insertion. Finally, the diazo compound in entry 4 presented the option of five- and six-membered ring formation from C-H insertion and seven-membered heterocycle formation from N-H insertion. Thus the kinetic preference for five-membered ring formation completely overwhelms any tendency for N-H insertion. However, four-membered rings are also directly accessible with rhodium(II) catalysts.33





Seebach<sup>34</sup> and McKervey<sup>35</sup> and their collaborators have found that Boc- and Cbz-protected  $\alpha$ -aminodiazoketones derived from L-alanine, L-valine, and L-phenylalanine cyclize very efficiently to the azetidin-3-one derivatives (eq. 11).



#### **1.5 Bisdiazo compounds**

The decomposition of bisdiazo compounds in rhodium(II) catalyst using in the synthesis is limited studied. Recently, the photochemistry of bisdiazo compounds was studied for the synthesis of highly strained compounds (eq. 12).<sup>36</sup>



Also, the intramolecular insertion reaction of bisdiazo compounds was established as a useful route to cyclic ethers, especially some medium-ring and large-ring compounds. Kulkowit and McKervey used  $Cu(II)(acac)_2$  in benzene to catalyze the intramolecular O-H insertion reaction of a bisdiazo ketone (eq. 13).<sup>37</sup>



#### **1.6 Diamino Dicarboxylic Acids**

Amino acid research is a large and important field. Diamino dicarboxylic acids or bis(amino acid)s are characterized by two glycine residues that are connected by a spacer via the  $\alpha$ -carbons. They contain two asymmetric carbons, two chemically identical amino



a bis(amino acid)

and carboxylic groups. This character confers on bis(amino acid)s an important role in studying biosynthesis inhibitors, such as investigating the synergistic effect when bacterial infections result in an immune response.<sup>38</sup> For example, bacterial peptidoglycan (Scheme 5) is a giant macromolecule consisting of linear heteroglycan chains cross-linked by short peptide chains. When studying the synergistic effect, pure muramyl peptides with different lengths of peptide chain are required. The bis(amino acid)s then

function as cross-linkers in the peptidoglycan network.



DAA = diamino acid or L-lysine (AA)<sub>n</sub> = n amino acids, with n = 0 - 5

#### Scheme 5

Also, the use of diamino dicarboxylic acid derivatives for peptidomimetic drug design is becoming more commonplace.<sup>39</sup> Recently, members of this class have been used as conformational constraints in order to mimic the secondary structures of peptides, such as  $\beta$ -turns,<sup>40</sup> and to stabilize a helical conformation.<sup>41</sup> As most peptides are flexible and have many conformations in solution which is difficult to determine the activity of the peptide. The conformational constraints may be helpful in elucidating these structures by restricting the peptide to a particular conformation or closely related family of conformations.<sup>42</sup> After being bridged or linked by a bis(amino acid), the peptide will be restricted to a particular conformation which will be associated with a receptor and a desired biological effect can be obtained (Scheme 6).<sup>45(a)</sup>



Scheme 6

## 1.7 Diaminoglutaric Acid and Its 3-Substituted Derivatives

Recently, much interest has been focused on the synthesis of unnatural and

unusual  $\alpha$ -amino acids,<sup>43</sup> since this class of compounds has an intrinsic biological activity. Such compounds can also modify biological potency and improve metabolic stability in a useful way when incorporated into medicinally important peptides.<sup>44</sup> Meanwhile, the synthesis of the modified peptides based on the substitution of  $\alpha$ -amino acids by non-standard residues in order to control flexibility and to determine conformations or to change the bioavailability and the inhibitory activity of the peptide has attracted significant attention.<sup>43(b),44,45</sup> In particular, the incorporation of  $\alpha$ -amino acids with a functional group that can act as a receptor ligand is of great interest.

#### **1.7.1 Diaminoglutaric Acid**

Glutaric acid is an important chemical messenger which is released in most of the excitatory synapses of mammalian central nervous systems.<sup>46</sup> Several syntheses of differentially functionalized (2R,4S)- and (2R,4R)-diaminoglutaric acids (DAG) **21** and **22** which was employed as a building block for peptidomimetics have been reported.<sup>47</sup>



Scheme  $7^{47(d)}$  and Scheme  $8^{47(f)}$  summarize two reported routes to **21** and **22**.



Scheme 7



The full syntheses are moderately lengthy. In work leading to his Ph. D. degree, Dr. Minmin Yang of our laboratory discovered that the N-H insertion reaction of diethyl 2-diazoglutaconate was accompanied by a shift of the double bond (Scheme 9). This

created the possibility for another diazotization/N-H insertion sequence. The net result was a concise high-yield synthesis of a bis(amino acid) derivative. When the final reduction was a hydrogenation over a chiral catalyst, partial control of stereochemistry was afforded.<sup>47(g)</sup>



Scheme 9

#### 1.7.2. 3-Substituted Diaminoglutaric Acids

The incorporation of  $\alpha$ -amino acids with a functional group that can act as a receptor ligand, such as 3-substituted diaminoglutaric acids, has attracted interest as possible peptidomimetics. Our target **27** here is to introduce fluorine into the  $\beta$ -position of diaminoglutaric acid.



It has been known for some time that fluorine can have profound and unexpected results on biological activity.<sup>48</sup> Once introduced, the high carbon-fluorine bond energy renders the substituent relatively resistant to metabolic transformations. The electronegativity of fluorine can have pronounced effects on the electron distribution in the molecule, affecting the basicity or acidity of neighboring groups, dipole moments within the molecule and the overall reactivity and stability of neighboring functional groups. As a consequence of the available electron density, fluorine can function as a hydrogen bond acceptor.<sup>49</sup> Introduction of fluorine can be carried out successfully in the presence of various functional groups using diethylaminosulfur trifluoride (DAST) as a fluorodehydroxylation reagent (Scheme 10).<sup>50</sup>



Scheme 10

We propose that the fluorine of 27 could be introduced this way (Scheme 11).



Scheme 11

The importance of enzymatic decarboxylations of amino acids in biosynthetic pathways suggested the utility of specific inhibitors of the decarboxylation enzymes in studying these pathways.<sup>49</sup> Fluoroacylated amino acids have been recognized as potent suicide inhibitors of enzymatic decarboxylation reactions.<sup>51</sup> The enzymatic inactivation is thought to be dependent upon loss of fluoride from the intermediate base **29** formed between pyridoxal phosphate and the fluoroacyl amino acid. Loss of fluoride generates a reactive acceptor **30** which can add an enzyme bound nucleophilic functional group. The covalently bound enzyme **31** is no longer free to bind additional substrate (Scheme 12). This character gives the fluoro-substituted amino acids high biological potential to serve as inhibitors to disrupt polyamine synthesis, with important implications for various anti-biological agents.<sup>49</sup>

A short and easy synthesis route to our target compound **27** starting from cheap and commercial available compound **32** was developed as briefly shown in Scheme 13. The novel aspect of this proposed route is the use of bisdiazo compound **33** to introduce both amino nitrogens simultaneously by double N-H insertion.


Scheme 12



Scheme 13

# 2. PART I. RESULTS AND DISCUSSION

#### 2.1. Synthesis of Bisdiazocarbonyl Compounds.

The bisdiazocarbonyl compound could be synthesized by diazo transfer from sulfonyl azide 7.<sup>52</sup> Dropwise addition of aqueous sodium azide to a stirred solution of *p*-acetoamidobenzenesulfonyl chloride **13** in methylene chloride containing tetrabutyl-ammonium bromide (TBAB) at 0 °C, gave *p*-acetoamidobenzenesulfonyl azide (*p*-ABSA) **7**,<sup>53</sup> as colorless crystals in 91.3% yield.



Compound **7** in acetonitrile was added dropwise to an acetonitrile solution of dimethyl 3-oxoglutarate **32** and triethylamine at 0 °C, in hopes of preparing **33** (eq. 14).



The resulting yellow oil showed only one methyl ester peak in its <sup>1</sup>H NMR spectrum (Figure 1). The peak at 74.2 ppm in the <sup>13</sup>C NMR spectrum (Figure 2), and the strong peak at 2133 cm<sup>-1</sup> in the IR spectrum (Figure 3) indicated that the C=N=N moiety was present in **33**. Other peaks in the <sup>13</sup>C NMR spectrum can be reasonably assigned as follows: 52.8 ppm, -CH<sub>3</sub>; 161.4 ppm, ester carbonyls; 174.5 ppm, ketone carbonyl.



Figure 1. <sup>1</sup>H spectrum of compound 33

A possible reaction product, **34**, is inconsistent with the IR results. It is also ruled out by the results of elemental analysis (calcd for  $C_7H_6N_4O_5$  (**33**) %C, 37.18; %H, 2.67; %N 24.77: found %C, 37.15; %H, 2.69; %N, 24.53).



Thus the diazo transfer reaction afforded the hitherto unknown compound dimethyl 2,4-bis(diazo)-3-oxoglutarate **33** in excellent yield. (eq 15).



Figure 2. <sup>13</sup>C spectrum of compound 33



Figure 3. IR spectrum of compound 33

# 2.2. N-H insertion of Rh(II)-stabilized carbenoid from bis(diazo) compound 33 into Cbz-NH<sub>2</sub>.

Benzyl carbamate (Cbz-NH<sub>2</sub>) **35**, which was required for the N-H insertion reaction of dimethyl 2,4-bis(diazo)-3-oxoglutarate **33**, was easily afforded by bubbling dry ammonia gas through a methylene chloride solution of benzyl chloroformate at 0 °C (eq 16).



Dimethyl 2,4-bis(diazo)-3-oxoglutarate **33** reacted with excess benzyl carbamate **35** in refluxing methylene chloride, catalyzed by  $Rh_2(OAc)_4$  (eq 17). After the reaction solution had refluxed for 1 h, the TLC showed bis(diazo) compound **33** was consumed.



However, the purification of products was not smooth. The TLC exhibited a spot with a long tail when using 1:1 hexane:ethyl acetate as eluent. Trying various solvent systems,

such as methylene chloride with ethyl acetate, chloroform with methanol, and methylene chloride with toluene, did not solve the separation problem. Fortunately, when the reaction mixture was passed though a silica gel chromatography column eluted with 1:1 hexane:ethyl acetate, collecting all fractions before a yellow fraction on the column gave a mixture which could be separated by TLC using 5:1 hexane:ethyl acetate. Finally, a colorless oil was afforded by column chromatography in 63% yield. However, the <sup>1</sup>H NMR spectra (Figure 4) showed four single peaks from 3.67~3.75 ppm which indicated four kinds of methyl groups of the methyl ester. Initially, we thought that the separated colorless oil was still a mixture. However the use of various solvent systems to attempt to separate this oil by TLC failed. The M+1 peak indicated on mass spectrum and elemental analysis of **36** were successful, which means the N-H insertion reaction occurred and the desired product was formed. Why did the NMR spectra show a mixture?



**Figure 4a.** Partial <sup>1</sup>H NMR spectrum of **36**: OCH<sub>3</sub> region.



**Figure 4c.** Partial <sup>1</sup>H NMR spectrum of **36**:  $7.0 \sim 8.2$  ppm region.

It is reasonable to assume that compound **36** is highly enolic (Scheme 14). In Scheme 14, the carbons are numbered only to distinguish one end of the molecule from the other. That is, the enol double bond might be between C2 and C3, which we denote with a suffix "**a**", or it might be between C3 and C4, which we denote with a suffix "**b**".



Furthermore, geometrical (i.e. cis-trans) isomerism is possible for these enols. Thus, as

Figure 5a. Partial <sup>13</sup>C NMR spectrum of 36: 52.5 ~ 54.6 ppm region. A DEPT-135

experiment showed all these peaks were from CH<sub>3</sub>/CH carbons.





experiment showed all these peaks were from CH<sub>2</sub> carbons.



Figure 5c. Partial  $^{13}$ C NMR spectrum of 36: 127.9 ~ 128.9 ppm region. A DEPT-135

experiment showed all these peaks were from CH<sub>3</sub>/CH carbons.



Figure 5d. Partial <sup>13</sup>C NMR spectrum of 36: 134.3 ~ 136.2 ppm region. A DEPT-135

experiment showed all these peaks were from quaternary carbons.



**Figure 5e.** Partial <sup>13</sup>C NMR spectrum of **36**: 151.1 ~ 170.3 ppm region. A DEPT-135 experiment showed all these peaks were from quaternary carbons.

shown in Scheme 14, we have four possible forms of **36**, namely *E*-**36a**, *E*-**36b**, *Z*-**36a**, and *Z*-**36b**. This explains the four methyl signals in the 3.67 ~ 3.75 ppm region of the <sup>1</sup>H NMR. In Figure 4a, two methyl signals are marked with ° and two with \*. The two ° peaks are equal in area, as are the two \* peaks. Since "**a**" forms and "**b**" forms are equal in energy,  $K_{eq} = 1$ , *i.e.*, there must be equal amounts of *E*-**36a** and *E*-**36b**, and equal





amounts of Z-36a and Z-36b. The  $a \leftrightarrows b$  equilibrium in this solvent (CDCl<sub>3</sub>) at ambient probe temperature is apparently slow on the 400 MHz NMR timescale, so separate signals for a and b forms are observed. However, there is no reason to expect *E* and *Z* isomers to be present in equal amounts. From integration of ° peaks versus \* peaks we find a 60:40 ratio. It is reasonable to speculate that the more abundant form is the *E* form, because the internal hydrogen bond in the *E* isomer appears stronger than that in the *Z* form, making the *E* form slightly more stable.

To help assign the remainder of the peaks in the <sup>1</sup>H and <sup>13</sup>C NMR spectra, model compound **37** was used.<sup>74</sup> It is shown in Figures 6 and 7 with *E*-**36**.



Figure 6. Model compound used to assign <sup>1</sup>H NMR chemical shifts in 36.

From Figure 6, we may predict the chemical shifts of the protons in compound **36**. The peaks can be reasonably assigned as follows (note: the peaks came out pairs indicate the two diastereoisomers, *E*-**36** and *Z*-**36**): 8.0 ppm (broad peaks), CbzNHC=C (H*a*); 5.8 ppm and 5.9 ppm, four peaks also indicate four kind of C=CCH (H*b*) as the methyl ester groups; 5.1 ppm, OCH<sub>2</sub>Ph (H*c*/H*c'*); 4.8 ppm and 4.9 ppm (broad peaks), CbzNHCH (H*d*); 7.2~7.5 ppm (multiple peaks), phenyl protons (H*e*/H*e'*); 3.6~3.7 ppm methyl groups (H*f*/H*f'*).

We may also assign the <sup>13</sup>C peaks of compound **36** regarding the model compound **37** (Figure 7) as follows: 54.5 and 54.7 ppm, 53.3 ppm and 53.2 ppm, methyl group (C*a* and C*a*'); other peaks around 53 ppm, HNCH (C*b*); 67.3~68.7 ppm, OCH<sub>2</sub>Ph (C*c* and C*c*'); 128~129 ppm, aromatic carbon (C*d* and C*d*'); 134.5 ppm and 134.7 ppm, NHC=COH (C*e*); 136.2 ppm and 136.4 ppm, NHC=COH (C*f*, due to the –OH, the

chemical shift is downfield); 151.6 ppm and 156.5 ppm,  $COOCH_2Ph$  (Cg and Cg'); 167.6 ppm and 167.8 ppm, CHCOOMe (Ch), 170.2 ppm and 170.5 ppm, C=CCOOMe (Ci).



Figure 7. Model compound used to assign <sup>13</sup>C NMR chemical shifts in 36.

# 2.3. N-H insertion of Rh(II)-stablilized carbenoid from bis(diazo) compound 33 into Boc-NH<sub>2</sub>.

The *t*-butyl carbamate (Boc-NH<sub>2</sub>) **40**, which was required for the N-H insertion reaction of dimethyl 2,4-bis(diazo)-3-oxoglutarate **33**, was easily afforded by bubbling dry ammonia gas through an ethanol solution of *t*-butyl dicarbonate **40** at 0 °C. (eq 18)

$$Boc_{2}O + NH_{3(g)} \xrightarrow{CH_{2}CI_{2}} Boc-NH_{2}$$
(18)  

$$40 \qquad 41,96\%$$
  

$$\left(Boc = t-BuO \xrightarrow{O} \xi\right)$$

Dimethyl 2,4-bis(diazo)-3-oxoglutarate **33** reacted with excess *t*-butyl carbamate **41** in refluxing methylene chloride, catalyzed by Rh<sub>2</sub>(OAc)<sub>4</sub>. After the reaction solution had refluxed for 1 h, the TLC showed bis(diazo) compound **33** was completely consumed. Further purification by silica gel chromatography provided a colorless oil in 46% yield. As discussed in section **2.2**, the N-H insertion reaction product **42** consists of two enol forms *E*-**42** and *Z*-**42**. The NMR spectrum indicated the ratio of these two isomers was E-42: Z-42 = 67:33.



Scheme 15

Finally, we were able to get bisprotected diamino dicarboxylic acids **36** and **42**. Stereocontrol of the insertion products was impossible because of extensive enolization. Nevertheless, the successful double N-H insertion reaction of the new bisdiazo compound **33** reflects a further synthesis value on X-H (X = C, N, O, S, Se, P, halogen) insertion reactions.

### 2.4. Catalytic hydrogenolysis of N-H insertion reaction products.

Reduction of the ketone functionality of **36** and **42** to an alcohol (or  $CH_2$ ) was the next goal. First we tried the reaction shown in eq. 19, hoping to reduce the central "ketone" while simultaneously deprotecting the amines by hydrogenolysis of the Cbz groups. We hoped that hydrogenation of these two enols **36** and **42** over Pd catalysts would achieve that goal. The reported hydrogenations of similar enolic substances caused us to be optimistic. (eq. 21).<sup>54</sup> The reaction was run in either methanol or 1,4-dioxane solvent.





R = Me, t-Bu, Ph R' = Et, Hmodifier - HCd or HCn to improve the *ee* HCd = 10,11-dihydrocinchonidine HCn = 10,11-dihydrocinchonine

After hydrogenation under 40 psi H<sub>2</sub>, compound **36** yielded white crystals after purification. However, the <sup>1</sup>H NMR spectrum gave an unexpected result. A doublet signal at 3.0 ppm coupled with triplet signal at 3.7 ppm indicated that there is a -CH<sub>2</sub>-CHgroup in the structure which did not match any groups of compound **28**. Fortunately, we were able to obtain the x-ray structure of this strange product **44** (Figure 8). The crystal structure data are included in Appendix A.



Figure 8. X-ray structure of compound 44

A big question came to our mind. Why would we get compound **44**? We examined a sample of **36** by NMR after establishing the structure of **44** to check whether **36** had decomposed or rearranged on storage. It had not.



After searching the literature, we found a few examples which seemed to be related to the mechanism by which **44** is produced. Equation 22 illustrates the unusual formation of N-benzylamines during Cbz group hydrogenolysis.<sup>67</sup> Equation 23<sup>75</sup> is an example which illustrates the formation of a nucleophilic intermediate (in this case an enolate) during catalytic hydrogenation. These precedents support us to propose a mechanism for the hydrogenolysis of compound **36** on Pd/C catalyst (Scheme 18).



$$(23)$$

We postulate that in several steps **49** is formed by routine deprotection of the "righthand" NHCbz group (as drawn in Scheme 16) and deprotection followed by C-N hydrogenolysis of the "lefthand" NHCbz group. The next step is similar to eq. 23 in which a nucleophilic species attacks a neighboring carbonyl carbon. Cyclopropane **50** opens to relieve ring strain, giving **51** and hydroxide. Hydroxide adds to **51** in Michael fashion to give **52**, which loses hydride (perhaps to **49**), giving **53**. This tautomerizes to **44**. This mechanism is highly speculative.



Scheme 16. Proposed mechanism for formation of 44

We also tried Pd/C catalyzed hydrogenation of Boc-protected compound **42** which won't undergo the hydrogenolysis of Boc group. However, we only found the starting compound **42** instead of expected alcohol **48**.



#### **3. PART II. INTRODUCTION**

#### **3.1 Introduction to hypervalency.**

The concept of hypervalent molecules was established by J. I. Musher in 1969.<sup>55</sup> Hypervalent compounds are ions or molecules of the elements bearing more electrons than the octet within a valence shell (Scheme 17). According to Musher,<sup>55</sup> there are essentially two ways to hold electrons beyond the octet within a valence shell: (1) make up a set of dsp<sup>3</sup> or d<sup>2</sup>sp<sup>3</sup> hybrid orbitals using higher-lying d orbitals or (2) make up highly ionic orbitals revising the basic idea of Lewis that a covalent bond is formed by a localized pair of two electrons.



Scheme 17. Some simple hypervalent molecules

Pimentel and Rundle,<sup>56</sup> in 1951, laid the basis for new development in this area by proposing the idea of the three-center four-electron (3c-4e) bond, employing the molecular orbital theory. Owing to progress in computing and the efforts of many scientists, the idea of the 3c-4e bond has become supported and is presently widely accepted.

The 3c-4e bond (Scheme 18) is an electron-rich, orbital-deficient bond and the nonbonding molecule orbital (NBMO) becomes HOMO while the antibonding orbital is the LUMO. Most hypervalent species will have a pseudo-trigonal bipyramidal geometry (TBP) that employs two types of bonding: hypervalent bonding to two apical ligands, and normal bonding to all equatorial ligands.<sup>57</sup>



Scheme 18. Three-center four-electron (3c-4e), hypervalent bonding scheme

# 3.2 Hypervalent boron 10-B-5 species.

During the period when the interest in hypervalent species grew, very few hypervalent compounds of second-row elements were isolated or even detected. Only boron, carbon and fluorine species have been made so far, and there are only a small number of examples of these compounds. Some examples are the first hypervalent 10-B-5 boron compound **54** reported by Lee and Martin in 1984,<sup>58</sup> hypervalent 10-C-5

carbon compound **55** synthesized by Kin-Ya Akiba recently<sup>59</sup> (Scheme 19) and first 10-F-2 hypervalent fluorine species observed by Ault and Andrews in 1976,<sup>60</sup> which was the highly unstable trifluoride anion  $F_3^-$  generated by simultaneous deposition of an argon/fluorine mixture with CsF, RbF or KF and detected at low temperature by Raman and infrared spectroscopy.



Scheme 19. Some hypervalent compounds based on second row elements.

Recently, Wada et al.<sup>61</sup> reported a synthesis route to tris(2,6-dimethoxyphenyl)borane **56** which let us set a program leading to a new 10-B-5 species **59** (Scheme 20). In this thesis, we report on progress toward this goal.



Scheme 20

# 3.3 Hypervalent phosphorus 10-P-5 species.

Phosphorus species **61** also attracted our interest (Scheme 23). It is formally an 8-P-5 species that may exhibit a three-center two-electron (3c-2e) bond. An analogous compound, **65**, was described recently.<sup>72</sup> Compound **65** was found to be undergoing a



dynamic equilibrium in solution. The x-ray crystal structure of **65** revealed an unsymmetrical B-N-B triad (*i.e.* 65b or 65c), with the "free" boron coordinated to a

solvent THF molecule. It was of interest to see whether **61**, the phosphorus analogue of **65**, would have a greater tendency than **65** to engage in three-center bonding.

Hui Li of our laboratory has made progress in the synthesis of **61**.<sup>73</sup> We report herein our work directed at the completion of this synthesis.



Scheme 21

#### 4. PART II. RESULTS AND DISCUSSION

#### 4.1 Synthesis of tris(2,6-dimethoxyphenyl)borane

Compound **56**,  $\Phi_3$ B (the 2,6-dimethoxyphenyl group is sometimes denoted as " $\Phi$ "), was prepared by the reaction of  $\Phi$ Li **58** and boron trifluoride diethyl etherate.<sup>61</sup> To a solution of 1,3-dimethoxybenzene **57** and a catalytic amount of *N*,*N*,*N'*,*N'*-tetramethyl-ethylenediamine (TMEDA) in diethyl ether was added a 2.5 M *n*-hexane solution of *n*-butyllithium at 0 °C under nitrogen. To the suspension of  $\Phi$ Li **58** after precipitates formed was added boron trifluoride diethyl etherate in benzene. After reflux and recrystallization from THF, white crystals of  $\Phi_3$ B **56** were obtained in 41% yield for both steps (eq. 24).



# 4.2 Demethylation of tris(2,6-dimethoxyphenyl)borane, 56.

Many reagents have been utilized for demethylation of aryl methyl ethers as reported in the chemical literature. The most commonly used are hydrogen iodide (or bromide),<sup>62</sup> boron tribromide,<sup>63</sup> trimethylsilyl iodide,<sup>64</sup> aluminum chloride,<sup>65</sup> pyridinium hydrochloride,<sup>65</sup> and sodium amide.<sup>66</sup> In our research here, we have tried aluminum trichloride and boron tribromide so far.

# 4.2.1 Demethylation of 56 by aluminum trichloride.

To the fine suspension of  $\Phi_3$ B **56** in toluene was added anhydrous AlCl<sub>3</sub> under nitrogen. After reflux, water was added to hydrolyze the excess AlCl<sub>3</sub>. However, the white solid obtained proved to be resorcinol **66** by NMR spectrum (eq. 25). The unexpected decomposition indicates that the carbon-boron bonds in **56** were cleaved, although the demethylation still occurred.



# 4.2.2 Demethylation by BBr<sub>3</sub><sup>•</sup>S(Me)<sub>2</sub>.

BBr<sub>3</sub> is another commonly used demethylation reagent which came to mind after AlCl<sub>3</sub> failed to afford desired tris(2,6-dihydroxyphenyl)borane. To the solution of  $\Phi_3$ B **56** in methylene chloride was added a solution of BBr<sub>3</sub>·S(Me)<sub>2</sub> complex in methylene chloride at 0 °C under nitrogen. The afforded mixture indicated that the demethylation only partially occurred and we still detect resorcinol in the product by NMR spectrum (eq. 26).



The demethylation of **56** to target product could not be accomplished by either AlCl<sub>3</sub> or BBr<sub>3</sub>. Because of the time limit, further investigation of the demethylation of **56** has not been done yet. When choosing other demethylation reagents, the cleavage of the carbon-boron bond of  $\Phi_3$ B **56** might be avoided by avoiding acidic conditions, which seem to promote C-B cleavage.

### 4.3 Synthesis of tris(2,6-dihydroxyphenyl)phosphonium chloride 63.

The synthesis of compound **63** has been well developed by Hui Li in his previous work in our laboratory<sup>73</sup> as shown in Scheme 21. Tris(2,6-dimethoxyphenyl)phosphine **64** can be prepared by adding the triphenyl phosphite to **58** which was the product of ortho-lithiation of 1,3-dimethoxybenzene **57** in THF. Further demethylation of tris(2,6-dimethoxyphenyl)phosphine **64** by anhydrous AlCl<sub>3</sub> in toluene afforded a red solid.





By recrystallization from EtOEt/MeOH, a single crystal suitable for x-ray diffraction was obtained. The crystal was found to be a co-crystal with the ethyl ether, which was revealed by proton NMR. This indicated the hydrogen bond between compound **63** and solvent is strong which also was found that compound **63** associated with ethyl acetate from Hui Li's research.<sup>73</sup> A structure diagram is provided in Figure 9 and crystal structure data are included in Appendix B. We are interested in the bond lengths and bond angles of phosphorus to the neighboring atoms which are indicated in Table 1.



Figure 9. Crystal structure of compound 63

	Bond	Bond	Torsion
Parameters	length (Å)	angle (°)	angle (°)
P(1)-C(13)	1.770(3)		
P(1)-C(7)	1.770(3)		
P(1)-C(1)	1.786(2)		
P(1)-H(1)	1.27(2)		
C(13)-P(1)-C(7)		114.84(12)	
C(13)-P(1)-C(1)		113.68(12)	
C(7)-P(1)-C(1)		112.37(12)	
H(1)-P(1)-C(1)		104.7(11)	
H(1)-P(1)-C(7)		104.0(10)	
H(1)-P(1)-C(13)		106.0(11)	
H(1)-P(1)-C(1)-C(2)			30.8(11)
H(1)-P(1)-C(7)-C(8)			20.5(11)
H(1)-P(1)-C(13)-C(14)			24.5(11)

 Table 1. Selected Structural Parameters of 63



Scheme 23

The P(1), C(1), C(7) and C(13) formed an asymmetric pyramid with the equal length of P(1)-C(7) and P(1)-C(13). A search of the crystallographic literature revealed several reports of structures of the type  $(Ar_3PH)^+$ . The majority of crystal structures which were found involved the triphenylphosphonium ion, Ph<sub>3</sub>PH<sup>+</sup>, **67**.<sup>70</sup> Other more relevant examples were: tris(2,4,6-trimethoxyphenyl)phosphonium tetrachloroiron(III) **68**<sup>68</sup>, bis{tris(2,4,6-trimethoxylphenyl)phosphonium} bis[( $\mu_2$ -chloro)-dichloroiron] **69**<sup>69</sup>, (2-hydroxyphenyl)diphenylphosphonium bromide **70**<sup>71</sup> (Scheme 25). Geometrical parameters of these phosphonium ions are compared to those of **63** in Table 2.

Parameters	63	<b>67</b> <sup><i>a</i></sup>	68	69	70
h (Å)	-	0.525(16)	0.481	0.541	0.542
average P-C bond length (Å)	1.771(8)	1.786(8)	1.783(6)	1.791(7)	1.786(2)
P-H bond length(Å)	1.27(2)	1.351(6)	1.438	1.612	1.413
average C-P-C bond angle (°)	113.6(2)	110.7(5)	113.0(3)	111.3(2)	111.2(2)
average C-P-H bond angle (°)	104.9(4)	108.2(14)	105.7(4)	107.7(6)	107.4(7)

<sup>*a*</sup> values are averages of all reported triphenylphosphonium ion structures

#### Table 2.

We first thought that the parameters of compound **63** would be similar to those of compound **70** because compound **70** has –OH on its phenyl ring. However, from the Table above, the P-C bond length, C-P-C bond angle and C-P-H bond angle of compound **63** are most close to compound **68**. The P-H bond length of compound **63** is much less than either the average P-H bond length of triphenylphosphonium ion or the other triarylphosphonium ions.

We have been able to synthesis tris(2,6-dihydroxyphenyl)phosphonium chloride **63** and determine its structure. The next step is to deprotonate **63** by base (Scheme 21) which is a key step in the synthesis of compound **61** and the study of multicenter, perhaps hypervalent, bonding between the phosphorus and boron or other main group elements.

#### 5. EXPERIMENTAL

**General:** Methylene chloride and acetonitrile were distilled over calcium hydride under nitrogen atmosphere. Reagents used in the syntheses were purchased from the Aldrich Chemical Company and were used without further treatment.

Melting points were recorded on a capillary melting point apparatus and are uncorrected. NMR spectra were obtained at room temperature from Bruker AV-400 spectrometer operating at 400 MHz for <sup>1</sup>H and 100 MHz for <sup>13</sup>C. TMS was used as internal standard. The results are presented as parts per million (ppm or  $\delta$ ) and coupling constants are reported in Hz.

Mass spectral (MS) data were obtained using either a Finnegan 3300 or a VG 7070E mass spectrometer using a solid probe. The results are presented in terms of intensity percentage relative to the base peak and probable fragmentation product.

Elemental analysis was performed by Atlantic Microanalytical lab, GA.

TLC was performed on Sorbent Technologies silica G TLC plates w/UV254. Preparative column chromatography employed Sorbent Technologies silica gel (60 Å, 32-63 μm).

*p*-Acetoamidobenzenesulfonyl azide,  $7.5^2$  To a suspension of *p*-acetamidobenzenesulfonyl chloride 13 (50.0 g, 0.214 mol) in 400 ml methylene chloride containing tetrabutylammonium bromide (0.200 g, 0.008 mmol) at 0 °C was added dropwise a solution of sodium azide (16.0 g, 0.248 mol, in 50 mL H<sub>2</sub>O). The reaction was allowed to warm up to rt and was kept stirring at rt for 16 h. The organic layer was separated and washed with water (2 x 75 mL) and saturated aqueous sodium chloride (2 x 50 mL), dried over Na<sub>2</sub>SO<sub>4</sub> and evaporated to give 46.9 g (91.3%) pure white solid, mp 107-110 °C. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 2.25 (s, 3H), 7.80 (m, 2H), 7.89 (m, 2H), 8.04(br, 1H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 25.0, 119.8, 129.2, 132.7, 144.2, 169.3.

**Dimethyl 2,4-bis(diazo)-3-oxoglutarate, 33.** To a stirred solution of dimethyl 3-oxoglutarate **34** (2.41 g, 13.9 mmol) and triethylamine (3.00 g, 29.7 mmol) in 45 mL acetonitrile at 0 °C was added dropwise *p*-acetamidobenzenesulfonyl azide 7 (7.20 g, 30.0 mmol) in 100 mL acetonitrile over 30 minutes. The reaction was allowed to warm to rt and stirred for 12 h. The reaction mixture was filtered and the filtrate was evaporated under reduced pressure. The residue was washed with ethyl acetate/hexanes (1:1), filtered, and the solvent was evaporated. Further purification by silica gel column chromatography (hexanes/ethyl acetate = 4:1) gave 2.86 g yellow oil (92%). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 3.83 (s). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 52.8, 74.2, 161.4, 174.5. IR (neat): 3004.6, 2956.0, 2133.3, 1719.0, 1601.2, 1433.9, 1321.4, 1192.3, 1126.2, 996.1, 750.1, 763.1, 698.9. Anal. Calcd. for C<sub>7</sub>H<sub>6</sub>O<sub>5</sub>N<sub>4</sub>: C, 37.18; H, 2.67; N, 24.77. Found: C, 37.15; H, 2.69; N, 24.53.

Benzyl carbamate, 35. To a stirred solution of benzyl chloroformate (10.3 g,

0.063 mol) in 200 mL methylene chloride was bubbled through dry ammonia gas at 0 °C. Stopped bubbling while there was no more precipitation formed. Filtered the precipitation and the filtrate was evaporated under reduced pressure to give 9.10 g (99%) white crystals, mp 87-88 °C. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 4.89 (br, 2H), 5.10 (s, 2H), 7.36 (m, 5H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 67.1, 128.3, 128.4, 128.8, 136.4, 157.0.

**Compound 36.** To a stirred solution of dimethyl 2,4-bis(diazo)-3-oxoglutarate 33 (0.300 g, 1.33 mmol) in 15 mL methylene chloride was mixed with benzyl carbamate 35 (0.401 g, 2.66 mmol). A catalytic amount of rhodium(II) acetate (24 mg, 0.05 mmol) was added. The reaction set up was then immediately moved to a pre-heated oil bath at  $42^{\circ}$ C. After refluxing for 3 h, the bis(diazo) compound was completely consumed. The reaction mixture was purified by silica gel column chromatography (first using hexanes/ethyl acetate = 1:1 washing out the fractions before yellow band. The afforded fractions were separated by hexanes/ethyl acetate = 5:1) gave 0.430 g (68%) colorless oil. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 3.67 (s, 3H), 3.71 (s, 3H), 4.96 (br, d, J = 3.2 Hz, 1H), 5.98 (d, J =10.3Hz, 1H), 7.95 (br, 1H) for one isomer; 3.73 (s, 3H), 3.75 (s, 3H), 4.79 (br, 1H), 5.82 (d, J = 9.2 Hz, 1H), 8.17 (br, d, J = 1.2 Hz, 1H) for the other isomer, 5.15 (m, 5H, two *CH*<sub>2</sub>Ph, O*H*), 7.33 (m, 10H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 52. 8, 53.2, 53.2, 53.3, 53.3, 53.4, 128.1, 128.3, 128.3, 128.4, 128.7, 128.7, 128.7, 128.8, 128.92, 128.94, 129.03, 129.03, 134.6, 134.7, 136.2, 136.4, 151.6, 156.2, 156.5, 167.6, 167.9, 170.2, 170.5.

t-Butyl carbamate, 41. To a 50 mL saturated NH<sub>3</sub>-EtOH solution was added

dropwise a solution of t-butyl dicarbonate **40** (21.3 g, 97.8 mmol) in 50 mL ethanol at 0 °C. While stirring, the solution was bubbled through dry ammonia gas. Stopped bubbling while there was no more precipitation formed. Filtered the precipitation and the filtrate was evaporated under reduced pressure. Added 50 mL hexane. Boiled at 65 °C for 30 minutes. Cooled down to 0 °C. Washed with hexane and dried to give 11.0 g (96%) white crystals, mp 106-108 °C. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 1.45 (s, 9H), 4.67 (br, 2H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 28.4, 79.8, 156.7.

**Compound 42.** To a stirred solution of dimethyl 2,4-bis(diazo)-3-oxoglutarate **33** (1.94 g, 8.58 mmol) in 100 mL methylene chloride was mixed with *t*-butyl carbamate **41** (2.10 g, 17.9 mmol). A catalytic amount of rhodium(II)acetate (40 mg, 0.1 mmol) was added. The reaction set-up was then immediately moved to pre-heated oil bath at 42 °C. After refluxing for 3 h, the bisdiazo compound was completely consumed. The reaction mixture was purified by silica gel column chromatography (first using hexanes/ethyl acetate = 1:1 washing out the fractions before yellow band. The afforded fractions were separated by hexanes/ethyl acetate = 5:1) gave 1.59 g (46%) colorless oil. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 1.44 (s, 9H), 1.48 (s, 9H), 3.74 (s, 3H), 3.75 (s, 3H), 4.90 (br, 1H), 5.06 (dd,  $J_A = 10.4$ Hz,  $J_B = 4.00$ Hz, 1H), 5.71 (d, J = 10.4Hz, 1H), 7.68 (br, 1H) for one isomer; 1.45 (s, 9H), 1.49 (s, 9H), 3.74 (s, 3H), 3.78 (s, 3H), 4.69 (br, 1H), 5.14 (dd,  $J_A = 9.1$  Hz,  $J_B = 3.6$  Hz, 1H), 5.55 (d, J = 9.2 Hz, 1H), 7.96 (br, 1H) for the other isomer. <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 28.1, 28.1, 28.4, 28.4, 52.8, 53.0, 53.1, 53.2, 54.7, 54.8, 62.3,

63.4, 80.3, 80.6, 83.6, 83.8, 128.4, 129.2, 150.0, 150.4, 155.6, 155.8, 168.0, 168.4, 170.8, 170.9.

**Tris**(2,6-dimethoxyphenyl)borane, 56.<sup>61</sup> To a solution of 1,3-dimethoxybenzene 57 (5.52 g, 40.0 mmol) and a catalytic amount of *N*,*N*,*N*',*N*'-tetramethylethylenediamine (TMEDA, 0.2 mL) in diethyl ether (50 mL) was added a 2.5 M n-hexane solution of *n*-butyllithium (16 mL, 40 mmol) at 0 °C under nitrogen. The mixture was stirred for 3 h to afford the precipitate of  $\Phi$ Li 58. To this suspension was added boron trifluoride diethyl etherate (1.89 g, 13.2 mmol) in benzene (50 mL) and the mixture was refluxed for 4 h. 15 mL methanol was added at rt and the insoluble materials were recrystallized from THF to give 2.27 g (41%) white crystals of  $\Phi_3$ B 56. m.p.: 234-235 °C. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 3.45 (s, 18H), 6.47 (d, *J* = 8.2 Hz, 6H), 7.18 (t, *J* = 8.2 Hz, 3H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 56.8, 105.3, 130.3, 162.7.

**Tris(2,6-dimethoxyphenyl)phosphine, 64.** To a solution of 1,3-dimethoxybenzene **57** (10.0 g, 72.5 mmol) in THF (40 mL) was added a 2.5 M *n*-hexane solution of *n*-butyllithium (30.5 mL, 76.3 mmol) at 0 °C under nitrogen. After stirring at 0 °C for 45 minutes, a solution of triphenyl phosphite (6.75 g, 21.8 mmol) in THF (20 mL) was added dropwise. Then the stirring was continued overnight at rt. The reaction was quenched by adding a few mL methanol. Filtered the solid, washed with ether, combined the filtrate and the ether. Evaporate the solvent and wash the solid by water. The solid was recrystallized from acetone/water to give 3.61 g (37.5%) white crystals. m.p.:
149-151 °C. <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>): 3.48 (s, 18H), 6.44 (dd,  $J_A = 8.2$  Hz,  $J_B = 2.9$  Hz, 6H), 7.13 (t, J = 8.0 Hz, 3H). <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>): 56.2, 104.5, 115.5, 128.7, 162.5, 162.6. <sup>31</sup>P NMR (162 MHz, CDCl<sub>3</sub>): -66.6.

**Tris(2,6-dihydroxyphenyl)phosphonium chloride 63.** Anhydrous AlCl<sub>3</sub> (810 mg, 6.07 mmol) was added to a fine suspension of tris(2,6-dimethoxyphenyl)phosphine **64** (442 mg, 1.00 mmol) in 15 mL dry toluene under nitrogen. The suspension was refluxed for 2 h and the stirred at rt overnight, which resulted in much gray-green precipitate. The reaction was quenched by slow addition of 15 mL 3 M HCl and the insoluble material changed color to pink. The stirring was continued until none of the precipitate stuck to the glass. The solid was collected by filtration and washed with diethyl ether, and dried on the vacuum line, which afforded 308 mg (80%) almost pure compound **63**. Further purification can be achieved by recrystallization from Et<sub>2</sub>O/MeOH, mp 234 °C. <sup>1</sup>H NMR (400 MHz, DMSO-*d*<sub>6</sub>): 6.36 (dd, *J* = 5.6 Hz, 7.6 Hz, 6H), 7.22 (t, *J*= 8.1 Hz, 3H), 8.37 (d, *J* = 534 Hz, 1H), 10.63 (s, 6H). <sup>13</sup>C NMR (100 MHz, DMSO-*d*<sub>6</sub>): 93.2 (d, *J*= 102 Hz), 107.0 (d, *J*= 5.7 Hz), 136.3, 162.3. <sup>31</sup>P NMR (162 MHz, DMSO-*d*<sub>6</sub>): -51.3 (d, *J* = 527 Hz).

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APPENDICES

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loop_	
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atom type scat dispersion real	
_atom_type_scat_dispersion_imag	
_atom_type_scat_source	
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'International Tables Vol C Tables 4.2	.6.8 and 6.1.1.4'
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International Tables Vol C Tables 4.2	.6.8 and 6.1.1.4'
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'x+1/2, -y+1/2, -z	
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cell length c	16 1160(15)
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	193(2)
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9366			
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-6			
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-21			
20			
2.20			
28.36			
2297			
2042			
>2sigma(I)			
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'SMART	SAINT'		
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'SMART, SAINT'
'SAINT'
'SHELXS-97 (Sheldrick, 1990)'
'SHELXL-97 (Sheldrick, 1997)'
'SHELXP-97 (Sheldrick, 1997)'
'SHELXCIF-97 (Sheldrick, 2000)'

### \_refine\_special\_details

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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  $F^2^> 2$  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.

Fsqd
full
calc
$(P)^{2}+0.0097P$ ] where $P=(Fo^{2}+2Fc^{2})/3'$
direct
difmap
geom
mixed
none
?

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_refine_ls_shift/su_mean	0.000

#### loop\_

\_atom\_site\_label atom site type symbol atom site fract x atom site fract y atom site fract z \_atom\_site\_U\_iso\_or\_equiv atom site adp type atom site occupancy \_atom\_site\_symmetry multiplicity atom site calc flag \_atom\_site refinement flags atom site disorder assembly atom site disorder group C4 C -0.0009(4) 0.59200(13) 0.81987(10) 0.0319(3) Uani 1 1 d ... O1 O 0.0052(3) 0.59020(10) 1.11748(7) 0.0391(3) Uani 1 1 d ... C1 C 0.1480(3) 0.61401(12) 1.05091(10) 0.0269(3) Uani 1 1 d ... C3 C 0.1634(3) 0.60757(13) 0.89644(10) 0.0279(3) Uani 1 1 d . . . H3A H 0.2089 0.6923 0.9030 0.033 Uiso 1 1 calc R . . H3B H 0.3299 0.5626 0.8903 0.033 Uiso 1 1 calc R . . O2 O 0.1083(3) 0.64643(14) 0.75549(7) 0.0491(4) Uani 1 1 d . . . C5 C -0.0324(6) 0.6382(2) 0.67762(12) 0.0630(6) Uani 1 1 d . . . H5A H 0.0646 0.6816 0.6347 0.094 Uiso 1 1 calc R . . H5B H -0.2088 0.6726 0.6840 0.094 Uiso 1 1 calc R ... H5C H -0.0484 0.5549 0.6614 0.094 Uiso 1 1 calc R ... C6 C 0.1031(4) 0.63393(17) 1.19590(11) 0.0452(4) Uani 1 1 d . . . H6A H -0.0179 0.6105 1.2404 0.068 Uiso 1 1 calc R ... H6B H 0.1153 0.7204 1.1938 0.068 Uiso 1 1 calc R ... H6C H 0.2782 0.6005 1.2066 0.068 Uiso 1 1 calc R ... O4 O 0.3522(3) 0.66760(13) 1.05216(8) 0.0478(4) Uani 1 1 d . . . C7 C 0.0245(3) 0.43010(11) 0.98115(9) 0.0224(3) Uani 1 1 d . . .

O3 O -0.2041(3) 0.53906(16) 0.81592(8) 0.0559(4) Uani 1 1 d . . . C2 C 0.0179(3) 0.56488(11) 0.97361(8) 0.0227(3) Uani 1 1 d . . . H2A H -0.1696 0.5922 0.9713 0.027 Uiso 1 1 calc R . . O5 O 0.24075(19) 0.37986(9) 0.98167(9) 0.0345(3) Uani 1 1 d . . . N1 N -0.2005(2) 0.37440(11) 0.98872(9) 0.0295(3) Uani 1 1 d . . . H1A H -0.2028 0.2970 0.9946 0.035 Uiso 1 1 calc R . . H1B H -0.3495 0.4144 0.9880 0.035 Uiso 1 1 calc R . .

loop\_

atom site aniso label atom site aniso U 11 atom site aniso U 22 atom site aniso U 33 atom site aniso U 23 atom site aniso U 13 atom site aniso U 12 C4 0.0338(8) 0.0317(7) 0.0301(7) 0.0003(5) 0.0030(7) 0.0076(7) O1 0.0437(7) 0.0430(6) 0.0305(6) -0.0058(4) 0.0011(5) -0.0132(6) C1 0.0281(7) 0.0195(6) 0.0330(7) 0.0015(5) -0.0038(6) -0.0002(5) C3 0.0262(7) 0.0250(7) 0.0325(7) 0.0032(6) 0.0022(6) -0.0012(6) O2 0.0593(9) 0.0562(8) 0.0319(6) 0.0092(6) 0.0019(6) -0.0044(7) C5 0.0833(17) 0.0782(14) 0.0275(9) 0.0073(9) -0.0005(9) 0.0111(14)  $C6\ 0.0591(12)\ 0.0450(10)\ 0.0316(8)\ -0.0056(7)\ -0.0023(8)\ -0.0069(9)$ O4 0.0435(8) 0.0570(8) 0.0430(7) -0.0002(6) -0.0051(6) -0.0242(7) C7 0.0199(6) 0.0199(5) 0.0275(6) 0.0005(5) -0.0005(6) 0.0006(5) O3 0.0470(8) 0.0825(11) 0.0381(7) 0.0015(7) -0.0084(6) -0.0213(8) C2 0.0187(6) 0.0199(5) 0.0294(7) 0.0005(5) -0.0006(6) 0.0003(5) O5 0.0185(5) 0.0220(5) 0.0630(7) 0.0038(5) 0.0009(5) 0.0016(4) N1 0.0196(6) 0.0207(6) 0.0481(8) 0.0020(5) -0.0001(5) 0.0001(4)

\_geom\_special\_details

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.

loop

\_geom\_bond\_atom\_site\_label\_1 \_geom\_bond\_atom\_site\_label\_2 \_geom\_bond\_distance \_geom\_bond\_site\_symmetry\_2 \_geom\_bond\_publ\_flag C4 O3 1.193(2) . ? C4 O2 1.327(2) . ? C4 C3 1.499(2) . ? O1 C1 1.322(2) . ? O1 C6 1.445(2) . ? C1 O4 1.199(2) . ? C1 C2 1.515(2) . ? C3 C2 1.524(2) . ? O2 C5 1.446(3) . ? C7 O5 1.2344(16) . ? C7 N1 1.3079(18) . ? C7 C2 1.5281(17) . ?

loop\_

geom angle atom site label 1 geom angle atom site label 2 \_geom\_angle\_atom\_site\_label\_3 geom angle \_geom\_angle\_site\_symmetry\_1 \_geom\_angle\_site\_symmetry 3 geom angle publ flag O3 C4 O2 123.42(16) . . ? O3 C4 C3 125.63(15) . . ? O2 C4 C3 110.95(15) . . ? C1 O1 C6 116.90(14) . . ? O4 C1 O1 124.17(14) . . ? O4 C1 C2 125.06(14) . . ? O1 C1 C2 110.77(12) . . ? C4 C3 C2 111.45(13) . . ? C4 O2 C5 116.30(18) . . ? O5 C7 N1 123.51(11) . . ? O5 C7 C2 118.57(12) . . ? N1 C7 C2 117.90(12) . . ? C1 C2 C3 110.14(12) . . ? C1 C2 C7 106.87(11) . . ? C3 C2 C7 111.70(12) . . ?

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0	
loop	
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atom type description	
atom type scat dispersion real	
atom type scat dispersion imag	
atom type scat source	
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\_symmetry\_cell\_setting

?

symmetry space group name H-M loop \_symmetry\_equiv\_pos\_as\_xyz 'x, y, z' '-x+1/2, y+1/2, -z+1/2' '-x, -y, -z' 'x-1/2, -y-1/2, z-1/2' cell length a 8.9250(10) cell length b 17.926(2) \_cell\_length\_c 14.692(2)cell angle alpha 90.000(2)cell angle beta 100.157(2)cell angle gamma 90.000(2)cell volume 2313.7(5) \_cell\_formula units Z 4 cell measurement temperature 193(2) \_cell\_measurement\_reflns used ? \_cell\_measurement theta min ? ? cell measurement theta max ? exptl crystal description ? exptl crystal colour \_exptl\_crystal\_size max ? ? exptl crystal size mid \_exptl\_crystal\_size min ? exptl crystal density meas ? exptl crystal density diffrn 1.346 \_exptl\_crystal\_density\_method 'not measured' exptl crystal F 000 984 \_exptl\_absorpt\_coefficient\_mu 0.274 exptl absorpt correction type ? exptl absorpt correction T min ? \_exptl\_absorpt\_correction\_T\_max ? exptl absorpt process details ? \_exptl\_special details ? ; \_diffrn\_ambient\_temperature 193(2) \_diffrn\_radiation wavelength 0.71073 \_diffrn\_radiation\_type MoK\a diffrn radiation source 'fine-focus sealed tube' 77

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_computing_publication_material	?

### \_refine\_special\_details

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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  $F^2^2 > 2$ 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.

\_refine\_ls\_structure\_factor\_coef Fsqd \_refine\_ls\_matrix\_type full \_refine\_ls\_weighting\_scheme calc \_refine\_ls\_weighting\_details

'calc w=1/[ $s^2(Fo^2)$ +(0.0831P)<sup>2</sup>+0.0000P] where P=(Fo<sup>2</sup>+2Fc<sup>2</sup>)/3' atom sites solution primary direct atom sites solution secondary difmap atom sites solution hydrogens geom refine ls hydrogen treatment mixed refine ls extinction method none refine ls extinction coef ? refine ls number reflns 5738 refine ls number parameters 288 refine ls number restraints 0 refine ls R factor all 0.1032 refine ls R factor gt 0.0587 refine ls wR factor ref 0.1575 refine ls wR factor gt 0.1436 refine ls goodness of fit ref 0.903 refine ls restrained S all 0.903 refine ls shift/su max 0.076 refine ls shift/su mean 0.009 loop atom site label atom site type symbol atom site fract x atom site fract y atom site fract z atom site U iso or equiv \_atom\_site adp type atom site occupancy atom site symmetry multiplicity atom site calc flag atom site refinement flags atom site disorder assembly atom site disorder group P1 P 0.65162(7) 0.08868(4) 0.25817(5) 0.03365(19) Uani 1 1 d ... Cl1 Cl 0.82487(8) 0.12525(4) 0.65688(5) 0.0484(2) Uani 1 1 d . . . O1 O 0.4273(2) -0.02282(10) 0.27741(15) 0.0565(6) Uani 1 1 d . . . H1A H 0.3615 -0.0546 0.2850 0.085 Uiso 1 1 calc R . . O2 O 0.4744(2) 0.22340(12) 0.17898(15) 0.0519(5) Uani 1 1 d . . . O3 O 0.8746(2) -0.02842(11) 0.28779(13) 0.0522(5) Uani 1 1 d ... H3A H 0.9348 -0.0648 0.2963 0.078 Uiso 1 1 calc R ... O4 O 0.5780(2) 0.11216(11) 0.04688(13) 0.0524(5) Uani 1 1 d . . . H4A H 0.6159 0.1522 0.0701 0.079 Uiso 1 1 calc R ... O5 O 0.7826(2) 0.20697(10) 0.14906(12) 0.0457(5) Uani 1 1 d . . . H5B H 0.8002 0.2452 0.1196 0.069 Uiso 1 1 calc R ... O6 O 0.7182(3) 0.13404(12) 0.44566(14) 0.0638(6) Uani 1 1 d . . . H6A H 0.7381 0.1468 0.5015 0.096 Uiso 1 1 calc R . .

O7 O 0.8237(3) 0.31142(13) 0.02966(16) 0.0811(8) Uani 1 1 d . . . C1 C 0.4502(3) 0.10155(14) 0.23108(17) 0.0336(6) Uani 1 1 d ... C2 C 0.3575(3) 0.04219(15) 0.24746(19) 0.0412(6) Uani 1 1 d . . . C3 C 0.2019(3) 0.05056(16) 0.2328(2) 0.0530(8) Uani 1 1 d . . . H3B H 0.1387 0.0102 0.2442 0.064 Uiso 1 1 calc R . . C4 C 0.1383(3) 0.11798(17) 0.2014(2) 0.0538(8) Uani 1 1 d ... H4B H 0.0308 0.1235 0.1917 0.065 Uiso 1 1 calc R . . C5 C 0.2254(3) 0.17673(16) 0.1841(2) 0.0452(7) Uani 1 1 d . . . H5A H 0.1792 0.2227 0.1625 0.054 Uiso 1 1 calc R . . C6 C 0.3814(3) 0.16895(14) 0.19835(18) 0.0366(6) Uani 1 1 d ... C7 C 0.7235(3) 0.04039(14) 0.17002(17) 0.0350(6) Uani 1 1 d . . . C8 C 0.8257(3) -0.01864(14) 0.1955(2) 0.0408(6) Uani 1 1 d . . . C9 C 0.8724(4) -0.06228(17) 0.1298(2) 0.0564(8) Uani 1 1 d . . . H9A H 0.9419 -0.1020 0.1474 0.068 Uiso 1 1 calc R ... C10 C 0.8187(4) -0.0484(2) 0.0384(2) 0.0697(10) Uani 1 1 d . . . H10A H 0.8494 -0.0801 -0.0069 0.084 Uiso 1 1 calc R ... C11 C 0.7212(4) 0.01013(19) 0.0097(2) 0.0590(9) Uani 1 1 d . . . H11A H 0.6870 0.0195 -0.0542 0.071 Uiso 1 1 calc R ... C12 C 0.6750(3) 0.05442(15) 0.07568(19) 0.0427(7) Uani 1 1 d . . . C13 C 0.7529(3) 0.17063(14) 0.29833(17) 0.0372(6) Uani 1 1 d ... C14 C 0.8077(3) 0.22225(14) 0.24043(18) 0.0389(6) Uani 1 1 d . . . C15 C 0.8841(3) 0.28535(16) 0.2765(2) 0.0519(8) Uani 1 1 d . . . H15A H 0.9213 0.3202 0.2371 0.062 Uiso 1 1 calc R . . C16 C 0.9059(4) 0.29722(18) 0.3701(2) 0.0621(9) Uani 1 1 d ... H16A H 0.9584 0.3408 0.3949 0.075 Uiso 1 1 calc R . . C17 C 0.8537(4) 0.24782(19) 0.4291(2) 0.0580(8) Uani 1 1 d . . . H17A H 0.8704 0.2572 0.4938 0.070 Uiso 1 1 calc R . . C18 C 0.7772(3) 0.18478(17) 0.39356(19) 0.0460(7) Uani 1 1 d . . . C19 C 0.8840(6) 0.2144(3) -0.0659(3) 0.123(2) Uani 1 1 d . . . H19A H 0.8843 0.2013 -0.1306 0.185 Uiso 1 1 calc R ... H19B H 0.8186 0.1796 -0.0395 0.185 Uiso 1 1 calc R ... H19C H 0.9880 0.2116 -0.0308 0.185 Uiso 1 1 calc R ... C20 C 0.8277(6) 0.2881(3) -0.0612(3) 0.1070(16) Uani 1 1 d . . . H20A H 0.8928 0.3228 -0.0893 0.128 Uiso 1 1 calc R ...

loop\_

atom site aniso label atom site aniso U 11 atom site aniso U 22 atom site aniso U 33 \_atom\_site\_aniso U 23 atom site aniso U 13 atom site aniso U 12 P1 0.0326(4) 0.0367(4) 0.0305(4) 0.0024(3) 0.0023(3) 0.0011(3) Cl1 0.0556(5) 0.0414(4) 0.0453(4) 0.0057(3) 0.0012(3) -0.0025(3) O1 0.0419(12) 0.0415(11) 0.0836(16) 0.0181(10) 0.0039(11) -0.0007(9) O2 0.0418(12) 0.0408(11) 0.0712(15) 0.0148(10) 0.0051(10) 0.0027(10) O3 0.0505(12) 0.0556(12) 0.0494(13) 0.0102(10) 0.0055(10) 0.0186(10) O4 0.0531(12) 0.0578(12) 0.0397(11) 0.0020(9) -0.0097(9) 0.0029(10) O5 0.0554(12) 0.0489(11) 0.0328(10) 0.0009(8) 0.0076(9) -0.0081(9) O6 0.0867(17) 0.0733(15) 0.0305(11) -0.0037(10) 0.0079(11) -0.0179(13) O7 0.143(2) 0.0575(14) 0.0476(14) 0.0038(11) 0.0293(15) -0.0024(15) C1 0.0291(13) 0.0396(14) 0.0313(13) -0.0001(11) 0.0034(10) -0.0001(11)  $C2\ 0.0372(15)\ 0.0430(15)\ 0.0425(16)\ 0.0030(12)\ 0.0049(12)\ 0.0039(12)$ C3 0.0363(16) 0.0519(18) 0.070(2) 0.0124(16) 0.0070(15) -0.0065(14) C4 0.0287(15) 0.065(2) 0.066(2) 0.0069(16) 0.0040(14) 0.0033(14) C5 0.0382(16) 0.0448(16) 0.0507(17) 0.0026(13) 0.0023(13) 0.0085(13)  $C6\ 0.0367(15)\ 0.0371(14)\ 0.0361(15)\ -0.0014(11)\ 0.0061(12)\ 0.0014(12)$  $C7\ 0.0323(14)\ 0.0380(14)\ 0.0339(14)\ -0.0025(11)\ 0.0036(11)\ -0.0038(11)$  $C8\ 0.0360(15)\ 0.0384(14)\ 0.0475(17)\ 0.0005(12)\ 0.0059(13)\ -0.0038(12)$ C9 0.057(2) 0.0498(18) 0.062(2) -0.0105(16) 0.0090(16) 0.0122(15)  $C10\ 0.080(2)\ 0.068(2)\ 0.062(2)\ -0.0288(19)\ 0.0163(19)\ 0.0061(19)$ C11 0.068(2) 0.070(2) 0.0361(17) -0.0135(15) 0.0017(15) -0.0012(17)  $C12\ 0.0402(16)\ 0.0461(16)\ 0.0398(16)\ -0.0035(13)\ 0.0015(12)\ -0.0048(13)$  $C13\ 0.0336(14)\ 0.0424(15)\ 0.0336(14)\ -0.0030(11)\ 0.0002(11)\ 0.0014(11)$  $C14\ 0.0337(14)\ 0.0454(15)\ 0.0361(15)\ -0.0017(12)\ 0.0019(12)\ 0.0021(12)$  $C15\ 0.0575(19)\ 0.0466(17)\ 0.0491(18)\ -0.0012(14)\ 0.0022(15)\ -0.0121(14)$  $C16\ 0.066(2)\ 0.0556(19)\ 0.057(2)\ -0.0094(16)\ -0.0087(17)\ -0.0156(17)$  $C17\ 0.066(2)\ 0.066(2)\ 0.0353(17)\ -0.0080(15)\ -0.0083(15)\ -0.0064(17)$ C18 0.0445(17) 0.0567(18) 0.0346(16) -0.0012(13) 0.0006(13) -0.0006(14)  $C19\ 0.190(6)\ 0.109(4)\ 0.087(3)\ 0.007(3)\ 0.069(4)\ 0.058(4)$  $C20\ 0.166(5)\ 0.103(4)\ 0.059(3)\ -0.002(2)\ 0.039(3)\ 0.012(3)$ C21 0.435(14) 0.100(4) 0.089(4) 0.018(3) 0.058(6) 0.115(7) C22 0.74(2) 0.151(7) 0.068(4) 0.010(4) 0.050(8) 0.196(11)

\_geom\_special\_details

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.

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loop geom bond atom site label 1 \_geom\_bond\_atom\_site\_label 2 geom bond distance \_geom\_bond\_site\_symmetry 2 geom bond publ flag P1 C13 1.770(3) . ? P1 C7 1.770(3) . ? P1 C1 1.786(2) . ? O1 C2 1.358(3) . ? O2 C6 1.344(3) . ? O3 C8 1.360(3) . ? O4 C12 1.368(3) . ? O5 C14 1.349(3) . ? O6 C18 1.354(3) . ? O7 C21 1.409(6) . ? O7 C20 1.405(4) . ? C1 C2 1.395(4).? C1 C6 1.401(3) . ? C2 C3 1.376(4).? C3 C4 1.379(4).? C4 C5 1.359(4).? C5 C6 1.379(4).? C7 C8 1.404(4) . ? C7 C12 1.400(4) . ? C8 C9 1.364(4) . ? C9 C10 1.367(5) . ? C10 C11 1.381(5).? C11 C12 1.371(4) . ? C13 C14 1.402(4) . ? C13 C18 1.401(4) . ? C14 C15 1.378(4).? C15 C16 1.371(4) . ? C16 C17 1.377(4) . ? C17 C18 1.375(4) . ? C19 C20 1.420(5) . ? C21 C22 1.075(8) . ?

loop

\_geom\_angle\_atom\_site\_label\_1 \_geom\_angle\_atom\_site\_label\_2 \_geom\_angle\_atom\_site\_label\_3

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_diffrn_measured_fraction_theta_max	0.995
_diffrn_reflns_theta_full	28.35
_diffrn_measured_fraction_theta_full	0.995
_refine_diff_density_max 0.465	
_refine_diff_density_min -0.400	
_refine_diff_density_rms 0.057	