



Article Optimization and a Kinetic Study on the Acidic Hydrolysis of Dialkyl α-Hydroxybenzylphosphonates

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Academic Editors: Milos Sedlak Received: 14 July 2020; Accepted: 17 August 2020; Published: 20 August 2020



Abstract: The two-step acidic hydrolysis of α -hydroxybenzylphosphonates and a few related derivatives was monitored in order to determine the kinetics and to map the reactivity of the differently substituted phosphonates in hydrolysis. Electron-withdrawing substituents increased the rate, while electron-releasing ones slowed down the reaction. Both hydrolysis steps were characterized by pseudo-first-order rate constants. The fission of the second P-O-C bond was found to be the rate-determining step.

Keywords: dialkyl α-hydroxyphosphonates; hydrolysis; phosphonic ester–acid intermediate; phosphonic acid; rate constants; mechanism

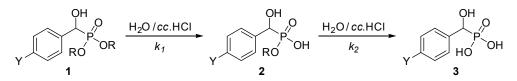
1. Introduction

The hydrolysis of P-esters (e.g., phosphinates and phosphonates) resulting in the formation of the corresponding acids (phosphinic acids and phosphonic acids, respectively) is an important chemical transformation, and hence it is applied widely in syntheses. Most often, the hydrolyses were performed under acidic conditions [1–4], but the application of NaOH or KOH is also common [5–8]. An additional possibility is the fission of the P-O-C unit by the effect of Me₃SiBr [9–11]. Usually, the acid- or base-catalyzed hydrolyses were carried out routinely, under "excessive" (unoptimized) conditions applying the acid or base catalysts in a larger quantity than required, and allowing longer reaction times. We undertook to explore the optimum conditions for the HCl-catalyzed hydrolysis of phosphinic and phosphonic esters. In the first round, the acid-catalyzed hydrolysis of cyclic phosphinates, such as 1-alkoxy-3-phospholene oxides, 1-alkoxyphospholane oxides, and an 1-alkoxy-1,2,3,4,5,6-hexahydrophosphinine oxide was investigated, optimized, and characterized by rate constants [12]. Then, the hydrolysis of a series of dialkyl arylphosphonates was studied. In this case, two-step conversions were monitored and quantified by k values [13]. α -Hydroxybenzylyphosphonates, obtained in the Pudovik reaction of substituted benzaldehydes and dialkyl phosphites, form a representative class of phosphonic acid derivatives [14]. α -Hydroxyphosphonates are versatile intermediates that may be transformed to α -aminophosphonates [15], can be phosphorylated [16], and may be rearranged to the corresponding phosphates [17]. The catalytic hydrogenation of α -dibenzyl hydroxyphosphonates afforded the respective α -hydroxyphosphonic acids [18]. Moreover, they may be of cytotoxic activity [18]. In this article, we describe our results on the HCl-promoted hydrolysis of α -hydroxyphosphonates and a few related analogues.

2. Results and Discussion

Substituted α -hydroxybenzylphosphonates (**1a**–**j**) prepared as described earlier [18] were subjected to acidic hydrolysis. The application of three equivalents (0.5 mL) of concentrated hydrochloric acid in

1 mL of water for ca. 2 mmol of the phosphonate (1) at reflux resulted in complete hydrolysis within 2.5–9.5 h depending on the substituents. The reactions followed a two-step protocol and took place via the corresponding ester–acid intermediate **2**, and were monitored by ³¹P NMR spectroscopy (Scheme 1). Experimental data together with the calculated pseudo-first-order k_1 and k_2 rate constants are listed in Table 1, while the concentration–time diagrams exhibiting the relative proportions of components **1**, **2**, and **3** are shown in Figures 1 and 2 and Figures S1–S8 in the Supplementary Materials.



Scheme 1. Two-step acidic hydrolysis of substituted α -hydroxybenzylphosphonates. For the Y substituents see Table 1.

Table 1. Experimental and kinetic data on the two-step hydrolysis of α -hydroxybenzylphosphonates **1a**–**j**.

Entry	Y	R	t _{max} (min)	t _r (h)	k ₁ (h ⁻¹)	k ₂ (h ⁻¹)	R ²
1	H (a)	Me	44	6.5	2.64	0.60	0.994
2	NO ₂ (b)	Me	22	2.5	5.18	1.24	0.989
3	Cl (c)	Me	34	5.5	3.36	0.79	0.987
4	F (d)	Me	32	6.0	3.93	0.67	0.965
5	CF ₃ (e)	Me	51	5.5	2.03	0.61	0.988
6	Me (f)	Me	76	8	1.64	0.31	0.962
7	H (g)	Et	90	9.5	1.03	0.35	0.986
8	$NO_2(h)$	Et	75	5.5	1.40	0.61	0.992
9	Cl (i)	Et	60	8.0	1.08	0.42	0.992
10	F (j)	Et	80	9.0	1.35	0.31	0.970

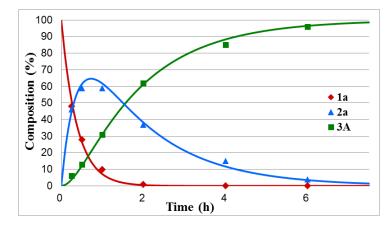


Figure 1. Concentration profile for the components during the hydrolysis of dimethyl α -hydroxybenzylphosphonate (1a) under optimum conditions. The R² measure of goodness of fit is 0.994.

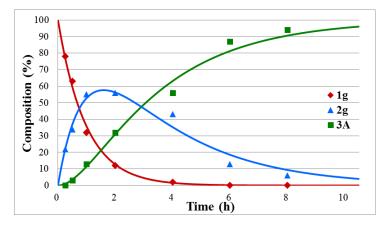


Figure 2. Concentration profile for the components during the hydrolysis of diethyl α -hydroxybenzylphosphonate (**1g**) under optimum conditions. The R² measure of goodness of fit is 0.986.

One can see that the hydrolysis of the unsubstituted dimethyl α -hydroxybenzylphosphonate was complete after $t_r = 6.5$ h, and the maximum proportion of intermediate **2a** could be observed at $t_{max} = 44$ min. In this case, k_1 and k_2 were found to be 2.64 h⁻¹ and 0.60 h⁻¹, respectively (Table 1/Entry 1). Electron-withdrawing substituents, such as 4-NO₂, 4-Cl and 4-F in the phenyl ring facilitated the hydrolyses that were complete after 2.5 h, 5.5 h, and 6 h, respectively. The maximum concentration of intermediates **2b–d** appeared in the range of 22–34 min. The k_1 values fell in the range of 3.36–5.18 h⁻¹, while the k_2 constants were between 0.67 and 1.24 h⁻¹ (Table 1/Entries 2–4). It seems that the 4-CF₃Ph substituent acted overall as the 4-ClPh group, as marked by $t_r = 5.5$ h. In this case (Y = CF₃), k_1 was 2.03, while k_2 was 0.61 (Table 1/Entry 5). In the above series, the hydrolysis of the 4-Me-substituted benzylphosphonate (**1f**) was the slowest, as a complete hydrolysis required 8 h, and the rate constants were found to be 1.64 (k_1) and 0.31 (k_2) (Table 1/Entry 6). It is noteworthy that the fission of the second P–OMe unit is the rate-determining step. As can be seen from Table 1, the k_1 values for the above cases are, in almost all cases, more than four times larger as compared to the k_2 values.

Regarding the series of substituted diethyl α -hydroxybenzylphosphonates (**1g**–**j**), the hydrolysis of the unsubstituted model (**1g**) was significantly slower than that of the dimethyl analogue (**1a**) (compare the reaction times of 9.5 h (Table 1/Entry 7) and 6.5 h (Table 1/Entry 1)). The corresponding k₁ and k₂ rate constants for the hydrolysis of diethyl ester **1g** were roughly the half the ones obtained for the methyl counterpart (**1a**) (compare rate constants 1.03/0.35 versus 2.64/0.60 (Table 1/Entry 7 versus Entry 1). Hydrolysis of the α -hydroxyphosphonates with electron-withdrawing 4-NO₂, 4-Cl, and 4-F substituents in the phenyl ring (**1h**-**j**) required shorter reaction times of 5.5–9.0 h as compared with that (9.5 h) of the unsubstituted instance (**1g**) (Table 1/Entries 8–10). In these cases again, the k₂ rate constants (0.61, 0.42, and 0.31, respectively) determined the overall reactivity.

To study the effect of substituents on the rate of the hydrolysis further, three additional model compounds, diethyl benzylphosphonate (**4k**), diethyl α -phenylethylphosphonate (**4l**), and diethyl β -phenylethylphosphonate (**4m**) were also subjected to hydrolysis, under the conditions applied for the α -hydroxybenzylphosphonates (**1a–j**) above (Scheme 2, Table 2, Figure 3 and Figures S9 and S10 in Supplementary Materials). It was found that the hydrolysis of the benzylphosphonate (**4k**) took longer than that of the α -hydroxy derivative **1g** (15 h versus 9.5 h, Table 2/Entry 1 and Table 1/Entry 7). The k₁ constant was somewhat larger for the hydrolysis of species **4k** than that for **1g**, but the decisive k₂ value become lower, as demonstrated by 1.12 h⁻¹ and 0.20 h⁻¹ versus 1.03 h⁻¹ and 0.35 h⁻¹ data pairs, respectively. Placing a Me group instead of the OH function on the α C atom, i.e., starting from α -phenylethylphosphonate **4l**, the hydrolysis became even slower, and it was complete only after 25 h. The smallest k values (k₁ = 0.51 h⁻¹, k₂ = 0.11 h⁻¹) were obtained in this case (Table 2/Entry 2). It is obvious that the lack of the electron-withdrawing OH group in position α , or the appearance of an

Me group instead of the HO function decreases the electrophilicity of the P atom of the P=O-function. The hydrolysis of β -phenylethylphosphonate (**4m**) with a reaction time of 20 h and k values of 0.70 h⁻¹ and 0.15 h⁻¹ (Table 2/Entry 3) occupied an intermediate position.

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Scheme 2. The two-step hydrolysis of other phosphonate derivatives. For the Y substituents see Table 2.

Entry	Y	t _{max} (h)	t _r (h)	k_1 (h ⁻¹)	k ₂ (h ⁻¹)	R ²
1	PhCH ₂ (k)	2	15	1.12	0.20	0.983
2	PhCHMe (1)	4	25	0.51	0.11	0.940
3	$Ph(CH_2)_2$ (m)	2.75	20	0.70	0.15	0.949

Table 2. Experimental and kinetic data on the two-step hydrolysis of phosphonates 4k-m.

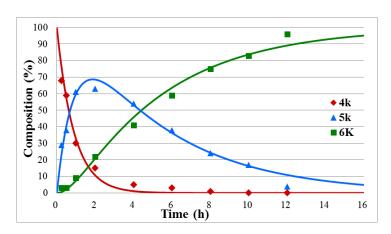


Figure 3. Concentration profile for the components during the hydrolysis of diethyl benzylphosphonate (**4k**) under optimum conditions. The \mathbb{R}^2 measure of goodness of fit is 0.983.

It is noted that the hydrolyses of the phosphonate function take place via the S_N^2 mechanism, i.e., by the nucleophilic attack of the water molecule on the P=O function. In the consecutive series, the fission of the second P-O-C bond is the rate-determining step.

The overall order of reactivity of the phosphonates (**1a**–**j** and **4k**–**m**) observed under acidic conditions was summarized in Table 3.

Table 3. Reactivity order of the phosphonates 1a-j and 4k-m in acidic hydrolyses characterized by t_r , as well as k_1 and k_2 .

	V Y MeO Y = NO ₂ , Cl, F	OH DMe > Ph MeO 1a	OMe ~ Etc Ih-j Y = NO ₂ , C	OEt Me	OH MeO OMe >	Ph PH O EtO OEt > PI	eto OEt > Ph.	$\frac{1}{10000000000000000000000000000000000$
t _r (h)	2.5-6.0	6.5	5.5–9.0	8	9.5	15	20	25
$k_1 (h^{-1})$	3.36-5.18	2.64	1.08 - 1.40	1.64	1.03	1.12	0.70	0.51
$k_2 (h^{-1})$	0.67–1.24	0.60	0.31-0.61	0.31	0.35	0.20	0.15	0.11

3. Materials and Methods

3.1. General Information

The ³¹P, ¹³C, and ¹H NMR spectra were taken on a Bruker DRX-500 spectrometer operating at 202.4, 125.7, and 500 MHz, respectively. The couplings are given in Hz. LC-MS measurements were performed with an Agilent 1200 liquid chromatography system coupled with a 6130 quadrupole mass spectrometer equipped with an ESI ion source (Agilent Technologies, Palo Alto, CA, USA). High-resolution mass spectrometric measurements were performed using a Thermo Velos Pro Orbitrap Elite hybrid mass spectrometer in positive electrospray mode.

3.2. Use of the ³¹P NMR Spectra in Quantitative Analysis

The composition of the reaction mixture was determined by the integration of the areas under the corresponding peaks of the starting material, intermediate, and product in the ³¹P NMR spectra.

3.3. Curve Fitting on the Time–Relative Quantity Data Pairs

The acidic hydrolysis was modeled assuming pseudo-first-order kinetics. The concentration of water and hydrochloric acid was constant during the reaction, and their initial concentration is incorporated in the pseudo-first-order rate constants k_1 and k_2 . The corresponding differential equations used in the model are the following:

$$\frac{d[\text{diester}]}{dt} = -k_1[\text{diester}] \tag{1}$$

$$\frac{d[\text{ester} - \text{acid}]}{dt} = k_1[\text{diester}] - k_2[\text{ester} - \text{acid}]$$
(2)

$$\frac{d[\text{acid}]}{dt} = k_2[\text{ester} - \text{acid}] \tag{3}$$

where [diester], [ester-acid], and [acid] are the time-dependent molarities of the dialkyl α -hydroxyphosphonate, the phosphonic ester–acid intermediate, and the phosphonic acid, respectively, and k_1 and k_2 are the pseudo-first-order rate constants of the first and the second step of the hydrolysis.

The solution of the differential equations is the following $(k_1 \neq k_2)$:

$$[\text{diester}] = c_0 e^{-k_1 t} \tag{4}$$

$$[\text{ester} - \text{acid}] = c_0 \left(\frac{k_1}{k_2 - k_1} e^{-k_1 t} - \frac{k_1}{k_2 - k_1} e^{-k_2 t} \right)$$
(5)

$$[\text{acid}] = c_0 \left(1 - \frac{k_2}{k_2 - k_1} e^{-k_1 t} + \frac{k_1}{k_2 - k_1} e^{-k_2 t} \right) = c_0 - [\text{diester}] - [\text{ester} - \text{acid}]$$
(6)

where c_0 is the initial molarity of the dialkyl α -hydroxyphosphonate.

The relative quantity of the components is their molarity divided by the sum of the three molarities (which is c_0). The calculated time–composition curves are described by the following equations, by leaving the k_1 and k_2 rate constants as parameters:

diester_{calcd} =
$$e^{-k_1 t} \times 100\%$$
 (7)

ester - acid_{calcd} =
$$\frac{k_1}{k_2 - k_1} \left(e^{-k_1 t} - e^{-k_2 t} \right) \times 100\%$$
 (8)

$$\operatorname{acid}_{calcd} = 100\% - \operatorname{diester}_{calcd} - \operatorname{ester} - \operatorname{acid}_{calcd}.$$
(9)

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During the calculation of the rate constants, we first gave arbitrary initial values to k_1 and k_2 , then, we optimized their values such that the sum of the squares of the differences between the experimental and calculated compositions became minimal:

$$SS_{res} = \sum_{i=1}^{n} \left(\text{diester}_{\exp,i} - \text{diester}_{\text{calcd},i} \right)^{2} + \sum_{i=1}^{n} \left(\text{ester} - \text{acid}_{\exp,i} - \text{ester} - \text{acid}_{\text{calcd},i} \right)^{2} + \sum_{i=1}^{n} \left(\text{acid}_{\exp,i} - \text{acid}_{\text{calcd},i} \right)^{2} \rightarrow \min$$

$$(10)$$

where *n* is the number of experimental time–composition data points measured at reaction times t_1 , t_2 , ..., t_n .

The resulting k_1 and k_2 rate constants and the associated time–composition curves were considered as the best fits. The best fits were found iteratively, using the nonlinear generalized reduced gradient method [19] of Microsoft Excel Solver.

The R² measure of goodness of fit is calculated as R² = $1 - \left(\frac{SS_{res}}{SS_{tot}}\right)$, where SS_{res} is described above and

$$SS_{tot} = \sum_{j=1}^{n} \left(\text{diester}_{\exp,j} - \frac{1}{n} \sum_{i=1}^{n} \text{diester}_{\exp,i} \right)^{2} \\ + \sum_{i=1}^{n} \left(\text{ester} - \text{acid}_{\exp,j} - \frac{1}{n} \sum_{i=1}^{n} \text{ester} - \text{acid}_{\exp,i} \right)^{2} \\ + \sum_{j=1}^{n} \left(\text{acid}_{\exp,j} - \frac{1}{n} \sum_{i=1}^{n} \text{acid}_{\exp,j} \right)^{2}$$
(11)

The reaction time (t_{max}) corresponding to the maximal ratio of the phosphonic ester–acid intermediate in the reaction mixture was found as follows:

$$0 = \left. \frac{d[\text{ester} - \text{acid}_{calcd}]}{dt} \right|_{t_{max}} = \left. \frac{d}{dt} \left(\frac{k_1}{k_2 - k_1} \left(e^{-k_1 t} - e^{-k_2 t} \right) \times 100\% \right) \right|_{t_{max}}$$
(12)

$$0 = \left. \frac{d}{dt} \left(e^{-k_1 t} - e^{-k_2 t} \right) \right|_{t_{max}} = -k_1 e^{-k_1 t_{max}} + k_2 e^{-k_2 t_{max}}$$
(13)

$$k_1 e^{-k_1 t_{max}} = k_2 e^{-k_2 t_{max}} \tag{14}$$

$$\frac{k_1}{k_2} = \frac{e^{-k_2 t_{max}}}{e^{-k_1 t_{max}}} \equiv e^{(k_1 - k_2) t_{max}}$$
(15)

$$t_{max} = \frac{\ln\left(\frac{k_1}{k_2}\right)}{k_1 - k_2}.$$
(16)

3.4. General Procedure for the Hydrolysis of Phosphonates (1a-j, 4k-m)

A mixture of 3.8 mmol of phosphonate (1a: 0.82 g, 1b: 0.99 g, 1c: 0.95 g, 1d: 0.89 g, 1e: 1.1 g, 1f: 0.87 g, 1g: 0.93 g, 1h: 1.1 g, 1i: 1.1 g, 1j: 1.0 g, 4k: 0.87 g, 4l: 0.92 g, 4m: 0.92 g), 1.0 mL (6.0 mmol) of *cc.* hydrochloric acid, and 2.0 mL of water was stirred at reflux for 2.5–25 h. The concentration of an aliquot part of the reaction mixture, or the whole mixture, afforded an oil that was analyzed by ³¹P NMR spectroscopy and LC-MS. Identification of the starting materials (1a–j, 4k–m), intermediates (2a–j, 5k–m), and products (3A–F, 6K–M) can be found in Table 4. The ¹³C and ¹H NMR spectral data of the new intermediates (2a–f, h–j) were obtained from the spectra of the corresponding mixtures containing also the phosphonic acids (3A–F).

	δ ³		
	Found (DMSO)	Literature	– [M + H]
1a	23.9	23.9 (CDCl ₃) [20]	216.9
1b	22.7	21.9 (CDCl ₃) [20]	262.2
1c	23.5	22.1 (CDCl ₃) [20]	273.0
1d	23.8	23.3 (CDCl ₃) [21]	235.0
1e [22] ^a	23.1	-	285.1
1f	24.1	24.0 (CDCl ₃) [20]	231
1g	21.8	21.5 (CDCl ₃) [20]	245.1
1h	20.4	19.9 (CDCl ₃) [20]	290.1
1i	21.2	21.9 (CDCl ₃) [20]	354.1
1j	21.7	21.7 (CDCl ₃) [23]	262.1
4 k	26.7	26.4 (CDCl ₃) [24]	229.2
41	29.8	30.4 (CDCl ₃) [25]	243.1
4m	30.6	30.8 (CDCl ₃) [24]	243.1
2a	20.8	19.9 (D ₂ O) [26]	203.04675 (203.04677 ^c)
2b ^b	18.8		248.03164 (248.03185 ^c)
2c ^b	20.3		237.00752 (237.00780 °)
2d ^b	20.4		221.03692 (221.03735 ^c)
2e ^b	19.6		271.03384 (271.03416 ^c)
2f ^b	20.8		217.06177 (217.06242 ^c)
2g [27] ^a	19.6		217.06193 (217.06242 ^c)
2h ^b	18.1		262.04707 (262.04750 ^c)
2i ^b	19.0		251.02306 (251.02345 ^c)
2j ^b	19.5		235.05267 (235.05300 ^c)
5k	23.7	25.5 (CD ₃ OD) [28]	201.1
51	27.0	32.3 (CDCl ₃) [29]	215.1
5m [30] ^a	27.5	_	215.1
3A	18.9	19.4(D ₂ O) [18]	189.1
3B	16.6	16.4 (D ₂ O) [18]	234.0
3C	17.9	18.6 (D ₂ O) [18]	223.0
3D	18.2	18.5 [31]	207.0
3E ^b	17.4		257.0
3F	18.6	18.8 (D ₂ O) [18]	203.1
6K	21.8	22.1 (DMSO) [32]	173.0
6L	25.7	27.8 (CD ₃ OD) [33]	187.0
6M	26.0	26.7 (DMSO) [34]	187.0

Table 4. Identification of the starting phosphonates (1a–j, 4k–m), ester–acid intermediates (2a–j, 5k–m), and phosphonic acids (3A–F, 6K–M).

^{a 13}C and ¹H NMR characterization is available in the literature; ^b Spectral data are given below; ^c Calculated value.

¹³C and ¹H NMR characterization of the new ester acids:

Methyl hydrogen 1-(4-*nitrophenyl*)-1-*hydroxymethylphosphonate* (**2b**). ¹³C NMR (DMSO-*d*₆) δ : 52.6 (d, ²*J* = 6.5, OCH₃), 69.1 (d, ¹*J* = 157.5, PCH), 122.7 (d, ⁴*J* = 2.4, C₃), 128.1 (d, ³*J* = 5.0, C₂), 146.5 (d, ⁵*J* = 3.3, C₄), 147.6 (C₁); ¹H NMR (DMSO *d*₆) δ : 3.58 (d, ³*J* = 10.4, 3H, OMe), 5.05 (d, ²*J* = 15.5, 1H, PCH), 7.64–7.73 (m, 2H, H₂), 8.17–8.23 (m, 2H, H₃).

Methyl hydrogen 1-(4-*chlorophenyl*)-1-*hydroxymethylphosphonate* (**2c**). ¹³C NMR (DMSO-*d*₆) δ : 53.0 (d, ²*J* = 6.5, OCH₃), 69.4 (d, ¹*J* = 160.3, PCH), 128.1 (d, ⁴*J* = 2.2, C₃), 129.5 (d, ³*J* = 5.3, C₂), 132.1 (d, ⁵*J* = 3.6, C₄), 139.0 (C₁); ¹H NMR (DMSO-*d*₆) δ : 3.55 (d, ³*J* = 10.3, 3H, OMe), 4.85 (d, ²*J* = 13.8, 1H, PCH), 7.32–7.47 (m, 4H, Ar).

Methyl hydrogen 1-(4-*fluorophenyl*)-1-*hydroxymethylphosphonate* (**2d**). ¹³C NMR (DMSO-*d*₆) δ : 53.0 (d, ²*J*_{P,C} = 6.5, OCH₃), 69.4 (d, ¹*J*_{P,C} = 161.3, PCH), 114.9 (dd, ²*J*_{F,C} = 21.2, ⁴*J*_{P,C} = 1.9, C₃), 129.7 (dd, ³*J*_{F,C} = 8.0, ³*J*_{P,C} = 5.6, C₂), 136.1 (d, ⁴*J*_{F,C} = 2.6), 161.9 (dd, ¹*J*_{F,C} = 242.5, ⁵*J*_{P,C} = 3.0, C₄); ¹H NMR (DMSO-*d*₆) δ : 3.54 (d, ³*J* = 10.2, 3H, OMe), 4.84 (d, ²*J* = 13.2, 1H, PCH), 7.09–7.19 (m, 2H, H₃), 7.38–7.49 (m, 2H, H₂).

Methyl hydrogen 1-(4-*trifluoromethylphenyl*)-1-*hydroxymethylphosphonate* (**2e**). ¹³C NMR (DMSO-*d*₆) δ : 53.1 (d, ²*J*_{P,C} = 6.5, OCH₃), 69.7 (d, ¹*J*_{P,C} = 157.9, PCH), 124.7–125.0 (m, C₃), 124.9 (q, ¹*J*_{F,C} = 271.1, CF₃), 127.3–128.7 (m, C₂, C₄), 145.0 (C₁); ¹H NMR (DMSO-*d*₆) δ : 3.56 (d, ³*J* = 10.3, 3H, OMe), 4.96 (d, ²*J* = 14.6, 1H, PCH), 7.57–7.72 (m, 4H, Ar).

Methyl hydrogen 1-(4-*methylphenyl*)-1-*hydroxymethylphosphonate* (**2f**). ¹³C NMR (DMSO-*d*₆) δ : 21.2 (Ar-CH₃), 52.8 (d, ²*J*_{P,C} = 6.6, OCH₃), 70.0 (d, ¹*J*_{P,C} = 160.2, PCH), 127.7 (d, ³*J* = 5.6, C₂), 128.7 (d, ⁴*J* = 2.1, C₃), 136.5 (d, ⁵*J* = 3.1, C₄), 136.9 (C₁); ¹H NMR (DMSO-*d*₆) δ : 2.28 (s, 3H, Ar-CH₃), 3.52 (d, ³*J* = 10.2, 3H, OMe), 4.76 (d, ²*J* = 13.0, 1H, PCH), 7.06–7.38 (m, 4H, Ar).

Ethyl hydrogen 1-(4-*nitrophenyl*)-1-*hydroxymethylphosphonate* (**2h**) ¹³C NMR (DMSO-*d*₆) δ : 16.9 (d, ³*J* = 5.6, CH₃), 62.0 (d, ²*J* = 6.5, OCH₂), 69.1 (d, ¹*J* = 157.5, PCH), 122.7 (d, ⁴*J* = 2.4, C₃), 128.2 (d, ³*J* = 4.9, C₂), 146.5 (d, ⁵*J* = 3.6, C₄), 147.6 (C₁); ¹H NMR (DMSO-*d*₆) δ : 1.17 (t, ³*J*_{H,H} = 7.0, 3H, CH₃), 3.91 (dq, ³*J*_{P,H} = 8.1, ³*J*_{H,H} = 7.0, 2H, OCH₂), 5.03 (d, ²*J* = 15.7, 1H, PCH), 7.65–7.72 (m, 2H, H₂), 8.16–8.23 (m, 2H, H₃). *Ethyl hydrogen* 1-(4-*chlorophenyl*)-1-*hydroxymethylphosphonate* (**2i**). ¹³C NMR (DMSO-*d*₆) δ : 16.6 (d, ³*J* = 5.5, CH₃), 61.7 (d, ²*J* = 6.5, OCH₂), 69.3 (d, ¹*J* = 160.7, PCH), 127.7 (d, ⁴*J* = 1.9, C₃), 129.2 (d, ³*J* = 5.3, C₂), 131.8 (d, ⁵*J* = 3.7, C₄), 138.6 (C₁); ¹H NMR (DMSO-*d*₆) δ : 1.15 (t, ³*J*_{H,H} = 7.1, 3H, CH₃), 3.91 (dq, ³*J*_{P,H} = 7.1, ³*J*_{H,H} = 7.1, 2H, OCH₂), 4.84 (d, ²*J* = 13.8, 1H, PCH), 7.33–7.47 (m, 4H, Ar).

Ethyl hydrogen 1-(4-*fluorophenyl*)-1-*hydroxymethylphosphonate* (**2j**). ¹³C NMR (DMSO-*d*₆) δ : 16.9 (d, ³*J* = 5.6, CH₃), 61.9 (d, ²*J* = 6.5, OCH₂), 69.5 (d, ¹*J*_{P,C} = 161.7, PCH), 114.8 (dd, ²*J*_{F,C} = 21.1, ⁴*J*_{P,C} = 2.0, C₃), 129.7 (dd, ³*J*_{F,C} = 8.2, ³*J*_{P,C} = 5.5, C₂), 136.1 (d, ⁴*J*_{F,C} = 2.8), 161.9 (dd, ¹*J*_{F,C} = 242.4, ⁵*J*_{P,C} = 3.2, C₄); ¹H NMR (DMSO-*d*₆) δ : 1.15 (t, ³*J*_{H,H} = 7.0, 3H, CH₃), 3.87–3.95 (m, 2H, OCH₂), 4.82 (d, ²*J* = 13.3, 1H, PCH), 7.11–7.17 (m, 2H, H₃), 7.42–7.47 (m, 2H, H₂).

1-(4-*Trifluoromethylphenyl*)-1-*hydroxymethylphosphonic acid* (**3e**). ¹³C NMR (DMSO-*d*₆) δ : 70.5 (d, ¹*J*_{P,C} = 157.7, PCH), 124.7–125.0 (m, C₃), 124.9 (q, ¹*J*_{F,C} = 272.3, CF₃), 127.2–128.6 (m, C₄), 128.4 (d, ³*J*_{P,C} = 5.0, C₂), 145.6 (C₁); ¹H NMR (DMSO-*d*₆) δ : 4.80 (d, ²*J* = 15.0, 1H, PCH), 7.55–7.72 (m, 4H, Ar).

4. Conclusions

Kinetic study of the two-step acidic hydrolysis of a series of dialkyl α -hydroxybenzylphosphonates and a few related model compounds allowed the mapping of the reactivity of the different substrates. The two-step hydrolyses were characterized by k_1 and k_2 pseudo-first-order rate constants belonging to the formation of the corresponding monoester monoacids and the phosphonic acids, respectively. Electron-withdrawing substituents increased the rate, while electron-releasing ones slowed down the hydrolyses starting with the nucleophilic attack of the water molecule. It turned out that the fission of the second P-O-C unit is the rate-determining step. The intermediate ester–acid species were identified and characterized.

Supplementary Materials: The following are available online: Figure S1: Concentration profile for the components during the hydrolysis of dimethyl α -hydroxy-4-nitrobenzylphosphonate (1b) under optimum conditions. The R² measure of goodness of fit is 0.989. Figure S2: Concentration profile for the components during the hydrolysis of dimethyl α -hydroxy-4-chlorobenzylphosphonate (1c) under optimum conditions. The R^2 measure of goodness of fit is 0.987. Figure S3: Concentration profile for the components during the hydrolysis of dimethyl α -hydroxy-4-fluorobenzylphosphonate (1d) under optimum conditions. The R² measure of goodness of fit is 0.965. Figure S4: Concentration profile for the components during the hydrolysis of dimethyl α -hydroxy-4-trifluoromethylbenzylphosphonate (1e) under optimum conditions. The R² measure of goodness of fit is 0.988. Figure S5: Concentration profile for the components during the hydrolysis of dimethyl α -hydroxy-4-methylbenzylphosphonate (1f) under optimum conditions. The R² measure of goodness of fit is 0.962. Figure S6: Concentration profile for the components during the hydrolysis of diethyl α -hydroxy-4-nitrobenzylphosphonate (1h) under optimum conditions. The R² measure of goodness of fit is 0.992. Figure S7: Concentration profile for the components during the hydrolysis of diethyl α -hydroxy-4-chlorobenzylphosphonate (1i) under optimum conditions. The R² measure of goodness of fit is 0.992. Figure S8: Concentration profile for the components during the hydrolysis of diethyl α -hydroxy-4-fluorobenzylphosphonate (1j) under optimum conditions. The R² measure of goodness of fit is 0.970. Figure S9: Concentration profile for the components during the hydrolysis of diethyl α -phenylethylphosphonate (41) under optimum conditions. The \mathbb{R}^2 measure of goodness of fit is 0.940. Figure S10: Concentration profile for the

components during the hydrolysis of diethyl β -phenylethylphosphonate (**4m**) under optimum conditions. The R² measure of goodness of fit is 0.949.

Author Contributions: Conceptualization, G.K. and Z.R.; methodology, N.H.; software, Á.S.; formal analysis, N.H. and J.K.; investigation, N.H.; resources, G.K.; data curation, Á.S./N.H.; writing—original draft preparation, G.K. and N.H.; writing—review and editing, G.K.; supervision, G.K.; project administration, G.K.; funding acquisition, G.K. All authors have read and agreed to the published version of the manuscript.

Funding: This project was funded by the National Research, Development and Innovation Office (K119202 and K134318).

Conflicts of Interest: The authors declare no conflict of interest.

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Sample Availability: Samples of the compounds are available from the authors.



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