


Cite this: *RSC Adv.*, 2020, **10**, 25898

# Phosphonopeptides containing free phosphonic groups: recent advances†

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Phosphonopeptides are mimetics of peptides in which phosphonic acid or related (phosphinic, phosphonous etc.) group replaces either carboxylic acid group present at C-terminus, is located in the peptidyl side chain, or phosphoramidate or phosphinic acid mimics peptide bond. Acting as inhibitors of key enzymes related to variable pathological states they display interesting and useful physiologic activities with potential applications in medicine and agriculture. Since the synthesis and biological properties of peptides containing C-terminal diaryl phosphonates and those with phosphonic fragment replacing peptide bond were comprehensively reviewed, this review concentrate on peptides holding free, unsubstituted phosphonic acid moiety. There are two groups of such mimetics: (i) peptides in which aminophosphonic acid is located at C-terminus of the peptide chain with most of them (including antibiotics isolated from bacteria and fungi) exhibiting antimicrobial activity; (ii) non-hydrolysable analogues of phosphonoamino acids, which are useful tools to study physiologic effects of phosphorylations.

Received 26th May 2020

Accepted 2nd July 2020

DOI: 10.1039/d0ra04655h

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## A Introduction

Phosphonopeptides can be broadly defined as mimetics of peptides in which carboxylic acid or peptide bond are replaced by phosphonic or related (phosphinic, phosphonous) moiety at any position of the peptide chain.<sup>1</sup> Also mimetics of

phosphopeptides with phosphate moiety being superseded by phosphonic acid may be included into that category.

On structural basis, phosphonopeptides could be divided into: (i) peptides, in which carboxylic acid group is replaced by phosphonic one (ii) peptides, in which phosphonic or phosphinic acid unit is mimicking peptide bond; and (iii) peptides in which phosphonic moiety replaces phosphate fragment in O-phosphonopeptides. Typical examples of the first class are: synthetic antibacterial agent alafosfalin,<sup>2</sup> phosphonic acid analogue of alanylalanine (compound 1) and natural antibiotic bialaphos,<sup>3,4</sup> a mimic of glutamylalanylalanine (compound 2),

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† This article is dedicated in honour of Professor Zbigniew Czarnocki on his 70th birthday.



*Paweł Kafarski was born in 1949. He studied chemistry at Wrocław University of Science and Technology where his scientific adventure started with M. Sc. Thesis, completed in 1971, followed by doctoral thesis (1977) both under the supervision of Prof. Przemysław Mastalerz. Prof. Mastalerz subsequently supervised his scientific career for many years. In his laboratory Paweł Kafarski worked on the synthesis of organophosphorus compounds and their potential biological activities. In 1976/1977 he interrupted his PhD studies and spent nine months at Marquette University at Milwaukee working in the laboratory of Prof. Sheldon E. Cremer on the synthesis of phosphetanes. In 1989 he spent six months in the laboratory of Prof. Henri-Jean Cristau at Ecole Nationale Supérieure de Chimie at Montpellier elaborating the procedure for the synthesis of phosphonopeptides containing P–N bond in their structures. Scientific activity of Paweł Kafarski was concentrated on elaboration of synthetic procedures suitable to produce phosphonate inhibitors (most likely in enantiomerically pure forms) of physiologically important enzymes, to mention only: aminopeptidases (targets for anti-cancer and anti-malarial drugs), cathepsin C (potential anti-tumor agents), glutamine synthetase (target for anti-tuberculosis agents), urease (antibacterials for treatment of stomach ulcer and stone formation in urinary tract) or L-phenylalanine ammonia lyase (potential herbicides). The design of ligands for these targets relied on knowledge of molecular mechanisms of the catalyzed reactions and on three-dimensional structures of the chosen proteins. He coauthored over 350 papers, which are well cited in the literature (over 6500 independent citations). He promoted 39 PhDs.*



both shown in Fig. 1. Diaryl (especially diphenyl) esters of such phosphonopeptides may be also classified to this group, however, will not be discussed in this review because synthesis and biological activity of these peptidyl diaryl phosphonates has been reviewed in detail.<sup>5–8</sup> Compound 3,<sup>9</sup> shown in Fig. 1 may serve as an example of this group. It acts as inhibitor of X-linked inhibitor of apoptosis protein thus being a potential anticancer agent. A potent inhibitor of several aminopeptidases – an analogue of homophenylalanyl-homophenylalanine (compound 4,<sup>10</sup> Fig. 1) represents the second group, whereas hexapeptide containing O-phosphotyrosine mimic (compound 5)<sup>11</sup> represents third group of phosphonopeptides. The latter one is one of the first effective inhibitors of protein-tyrosine phosphatase 1B, and thus became promising lead compound for treatment of type 2 diabetes. Also peptides containing phosphonate or phosphinate moiety in place of peptide bond will not be considered here since the synthesis and physiologic activities of these compounds are comprehensively reviewed.<sup>9,12–16</sup>

In this paper recent advances of chemistry and biology of phosphonopeptides holding free phosphonate or phosphinate groups will be discussed in some detail.

## B Peptides containing C-terminal aminophosphonic acids

Synthesis of peptides containing C-terminal phosphonic acid analogues of amino acids was intensively studied at the end of XX century. This was stimulated by promising activity and determination of molecular mode of action of alafosfalin (1) and bialaphos (2). Peptidyl parts of these molecules function as targeting unit. Thus, the peptides are efficiently transported through bacterial (or fungal) membranes and after hydrolysis release aminophosphonic acids, which exert their toxic action by inhibiting vital enzymes. In the case of alafosfalin it is an

analogue of alanine, a potent inhibitor of alanine racemase<sup>17</sup> and in the case of bialaphos it is phosphinothricin (compound 6), an inhibitor of glutamine synthetase and potent herbicide (Fig. 2).<sup>18</sup> This mechanism of action was found for majority of the phosphonopeptides isolated from bacteria and fungi (Fig. 3), namely: phosalacine (compound 7, releases phosphinothricin),<sup>19</sup> rhizocticins and plumbemycons (compounds 8, release inhibitor of threonine synthase),<sup>20</sup> and dehydrophos (compound 9).<sup>21</sup> The action of the latter one has quite unusual since phosphonic analogue of dehydroalaninate formed after cleavage of peptide bond is spontaneously converted into methyl acetylphosphonate (an analogue of pyruvic acid), which is strongly antibacterial by acting most likely as antimetabolite of pyruvate.<sup>22</sup>

The peptidyl structure of these antibiotics cause that they have limited utility in human medicine. It is because peptides are readily hydrolysed in human body fluids and thus obtained aminophosphonic acids are not able to cross bacterial cell barrier and exert antibiotic action.

Published in 2015 work of Metcalf and van der Donk brought a significant breakthrough in search for phosphonate antibiotics. By clever combination of the mining of genome of 10 000 of actinomycetes and selective labeling of phosphonate metabolites they rediscovered a huge number of old phosphonates and found out 19 new compounds including two phosphonopeptides – argolaphases (compounds 10, Fig. 3).<sup>23</sup>

A separate group of naturally occurring phosphonic peptidomimetics is the K-26 (compound 11)<sup>24</sup> and its analogues (Fig. 3), a family of bacterial secondary metabolites, tripeptides terminated by an unusual phosphonate analog of tyrosine. They are produced by three different actinomycetales and act as potent inhibitors of human angiotensin-I converting enzyme selectively targeting the eukaryotic family of the enzyme.<sup>25</sup>

Intensive search for synthetic bacterial phosphonopeptides was stimulated by the premise that phosphonic acid moiety, similarly to sulfonic one, may mimic carboxylic function. Consequently, aminophosphonic acids should act as false substrates for enzymes involved in amino acid metabolism and thus function as their inhibitors. Intensive studies on their inhibitory action towards various enzymes have indicated that this concept, in majority of cases, is invalid. Anyway, this

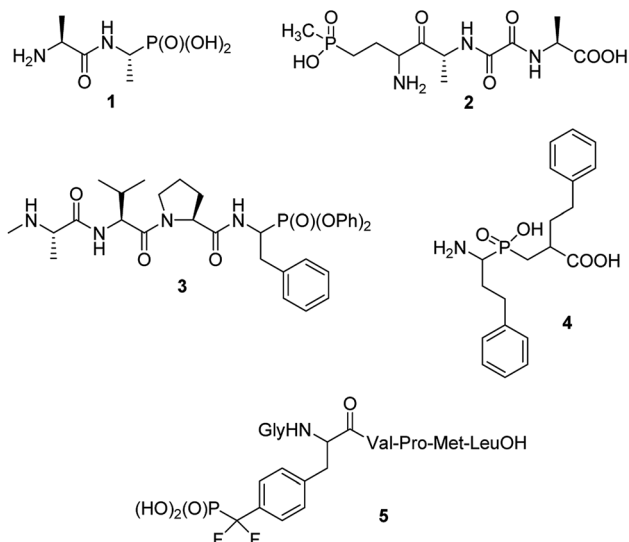


Fig. 1 Representative examples of structurally different phosphonopeptides.

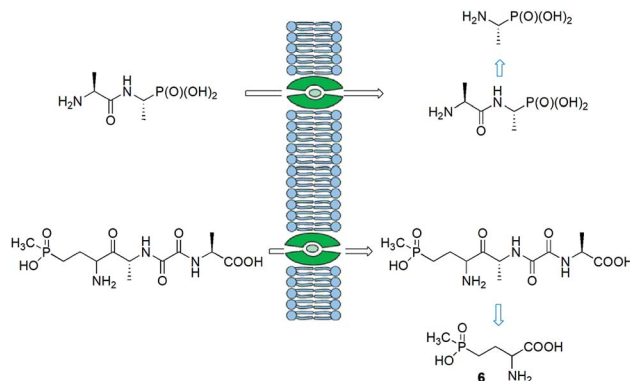


Fig. 2 Mechanism of action of alafosfalin and bialaphos.

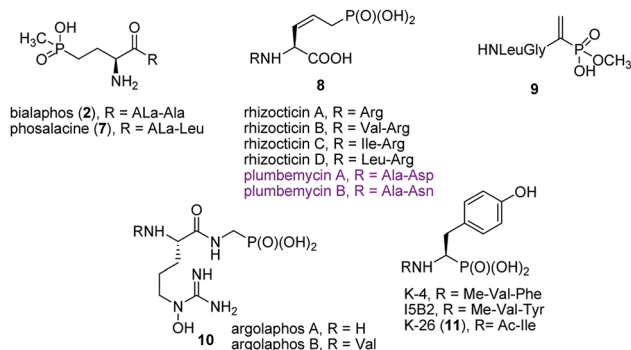


Fig. 3 Naturally occurring phosphonopeptide antibiotics.

assumption resulted in elaboration of effective procedures for preparation of this class of phosphonopeptides.<sup>26</sup>

Consequently, in last decade only singular papers devoted to the synthesis of phosphonopeptides, which used dialkyl aminophosphonates as substrates and applied classic peptide bond formation methods, have been published. They concern (see Fig. 4): synthesis of analogue of enkephalin (compound 12),<sup>27</sup> analgesic activity of which has not been described, N-substituted analogues of 1-(N-glycylamino)-1-1-methylethylphosphonic acids (compound 13 as representative example),<sup>28</sup> peptides of trifluoromethylated  $\beta$ -amino-phosphonates (compounds 14)<sup>29</sup> and phosphonic acid analogue of glutamic acid acylated with various enantiomers of prolyl-proline (for example, compound 15).<sup>30</sup> The latter peptides found an application in organic synthesis as effective bifunctional organocatalysts for enantioselective 1,4-Michael addition of aldehydes to nitroalkenes.<sup>31</sup>

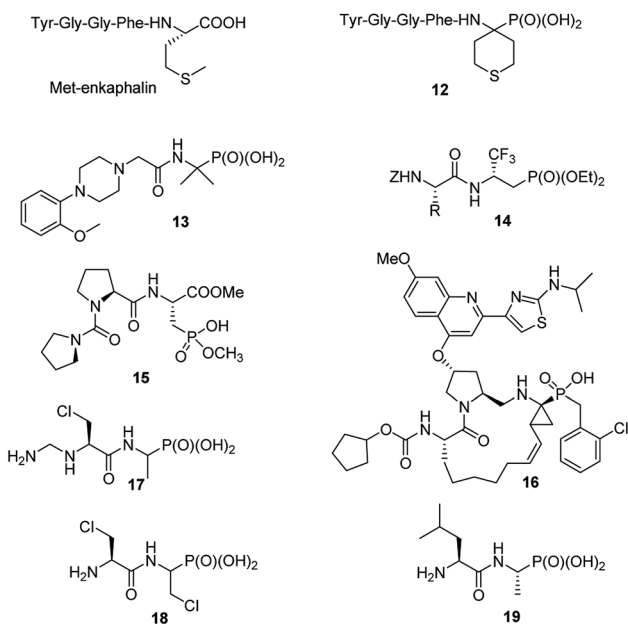


Fig. 4 Structurally variable peptides containing C-terminal amino-phosphonic acids.

Also classic liquid-phase peptide synthesis was applied for the preparation of a series of phosphonopeptides of quite complex structures being potent inhibitors of hepatitis C virus NS3 protease, with compound 16 (Fig. 4) showing the highest affinity to the enzyme. This compound also significantly inhibited proliferation of the virus in cell base assay.<sup>32</sup>

The resistance of bacteria to antibacterial agents is rising to dangerously high levels in all parts of the world. Simultaneously limited number of effective new antibiotics has been developed in the last few decades. One of the solutions here is to reinvestigate of old or abandoned drugs for reuse in medicine or for serving as lead compounds for the design of novel antibacterials.

Recently alafosfalin and its analogues, containing additionally second antibacterial agent -  $\beta$ -chloroalanine in the peptide chain, were synthesized by standard mixed carboxylic-carbonic anhydride method and were studied as new weapons against bacterial infections (for representative example see compounds 17 and 18 in Fig. 4).<sup>33,34</sup> Since the introduction of non-proteinogenic amino acids into peptide chain usually inhibits or slows its hydrolysis these compounds have a chance to be more stable in body fluids.

C-terminal domain of the dehydrophos (compound 9) biosynthetic enzyme DphH catalyzes the condensation of Leu-tRNA with phosphonic analogue of alanine, yielding dipeptide 19 (Fig. 4) – phosphonic acid analogue of leucylalanine. This enzyme is highly tolerant to structural variations in both aminoacyl-tRNAs and aminophosphonic acids and was used for the preparation of series phosphonodipeptides of interesting antibacterial properties.<sup>35</sup>

Quite original, approach was used to the synthesis of natural phosphono peptide – dehydrophos (Fig. 5).<sup>36</sup> Reaction started from readily available aminomethylenebisphosphonic acid, which was N-protected with carbobenzoxy (Z) moiety and then esterified by action of triethyl orthoformate. Removal of Z protection by hydrogenolysis followed by acylation with Boc-L-leucine resulted in peptide 20. This peptide was used as substrate for Horner–Wadsworth–Emmons reaction (a version of popular Wittig reaction), which gave dipeptide 21 containing C-terminal phosphonic analogue of dehydroalanine. Hydrolysis of its methyl group by action of strong base (Dabco®) followed by acidification with Dowex® is accompanied by rearrangement and decomposition of the desired dipeptide. Standard removal of Boc from compound 20 with trifluoroacetic acid followed by acylation with Fmoc-Gly an then Horner–Wadsworth–Emmons reaction resulted in peptide 22, which gave clean hydrolysis of one methyl group and after deprotection of Fmoc with Amberlite® afforded dehydrophos in nearly quantitative yield.

The limited number of papers dealing with application of classic methods of phosphonopeptide synthesis is not surprising if considering that they require accessibility to appropriate esters of aminophosphonic acids with free unblocked amino moiety, which have to be synthesized by independent procedures and are not readily available.<sup>37,38</sup>

Therefore, more efforts have been devoted to the elaboration of non-classical methods for preparation of peptides containing C-terminal aminophosphonic acids. One of the most obvious



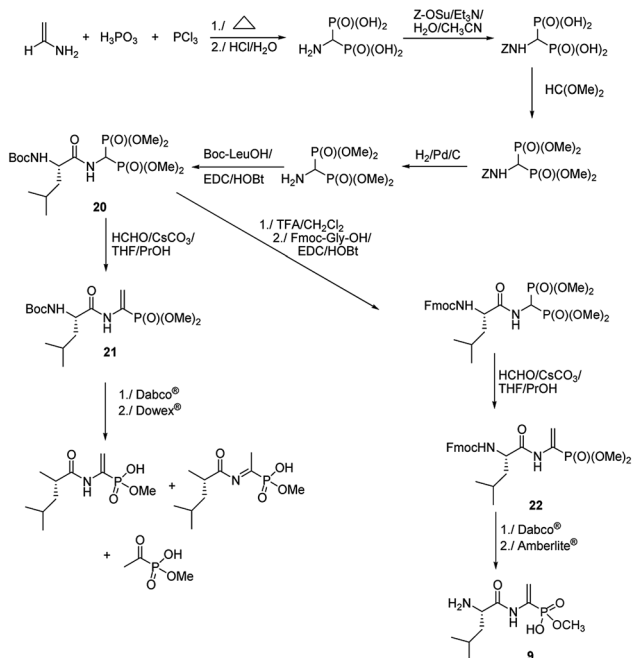


Fig. 5 Synthesis of dehydrophos.

solutions seems to be the introduction of phosphonate moiety at final steps of peptidomimetic synthesis. Thus, Mannich-type reaction of N-protected amides of amino acids and phosphorus trichloride (classical procedure for the preparation of aminophosphonic acids) or arylldichlorophosphine afforded nearly equimolar mixtures of diastereomers of the desired phosphonopeptides **23** with good yields (Fig. 6).<sup>39–41</sup> Similar reaction of N-protected taurine amides, aldehydes and arylldichlorophosphines was used to prepare sulphonophosphinodipeptides (compounds **24**, Fig. 6).<sup>41–43</sup>

Another approach is to use peptidylamides as substrates is their reaction with  $\alpha$ -isocyanatophosphonate, which afforded phosphono peptides with thiourea fragment replacing peptide bond (Fig. 7).

These peptides exhibited promising antiproliferative activity against several cancer cell lines (BCG-823, A-549, PC-3, Bcap-37 and BCG-823) with compounds **25**, **26**, **27** and **28** being the most active.<sup>44,45</sup>

An addition of diethyl or dimethyl phosphites to aldehydes derived from *S*-phenylalanyl-*S*-alanine resulted in diastereomeric mixtures of dipeptidyl 2-amino-1-hydroxyphosphonates

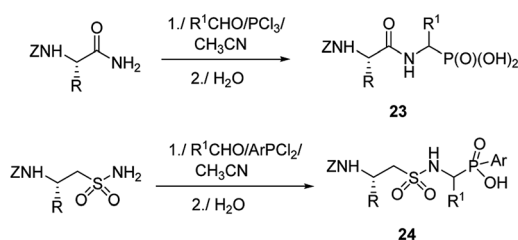


Fig. 6 Synthesis of phosphonopeptides by Mannich-like approach.

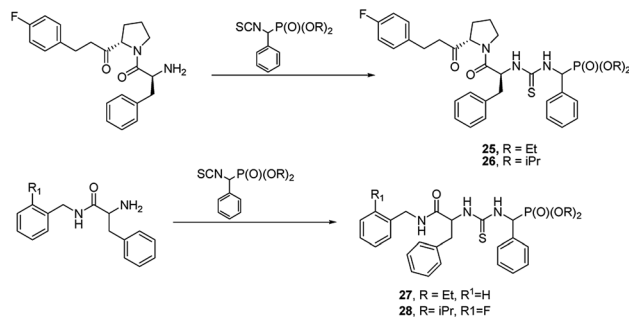


Fig. 7 Thiourea containing phosphonopeptides.

(Fig. 8). The formation of compounds of (SSS) configuration was privileged (72–92% de).<sup>46</sup> The obtained phosphonopeptides appeared to be promising inhibitors of cathepsins, with compound **29** (Fig. 9) being the most effective low-molecular weight inhibitor of cathepsin K described so far ( $K_i = 0.32$  nM).

Molecular modeling of binding of the most effective inhibitors towards cathepsins C and K suggest that hydroxyphosphonate fragment (but not phosphonic group as expected) acts as transition state analogue in this case (Fig. 9).<sup>46</sup> However, the possibility that hydroxyphosphonates decompose upon binding with the enzyme and release aldehyde, which is a real inhibitor, cannot be also ruled out.

An interesting example of synthesis of phosphonopeptides using aminophosphonic acids as substrates is an application of one-pot microwave-assisted multicomponent Ugi reaction. It provided a series of dipeptides **30** containing  $\alpha$ ,  $\beta$  and  $\gamma$  aminophosphonates as C-terminal ones (Fig. 10). A high level of structural diversity may be obtained by this approach providing a platform for the synthesis of novel peptidomimetics.

This reaction was also used to generate zwitterionic, chromatographic selectors (Fig. 10), which might be considered as new reversed-phase/ion-exchangers with interesting properties.<sup>47,48</sup>

Functionalization of the aminophosphonate of already existing peptides also seems to be an attractive approach for synthesis of more complex structures. An example of such an approach is a simple conversion of *N*-(1,2,2,2-tetrachloroethyl) amides (compounds **31**, Fig. 11), which were obtained from *N*-phthalylamino acid amides, into 1-acylamino-2,2,2-

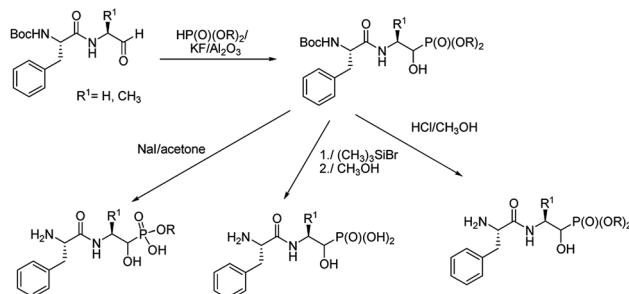


Fig. 8 Synthesis of peptides containing 1-hydroxy-2-aminoalkylphosphonates.



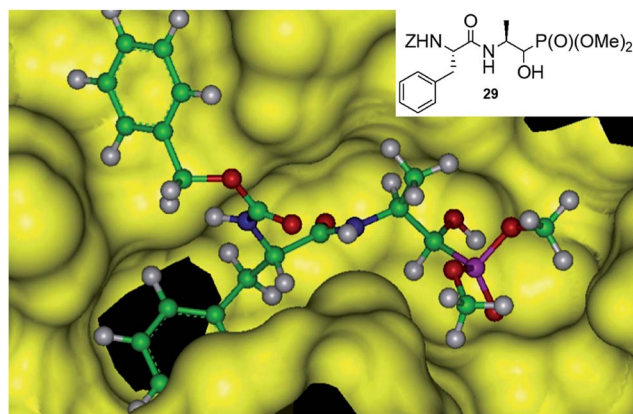


Fig. 9 Mode of binding of peptide 28 by cathepsin K.

trichloroethylphosphonates (compounds 32, Fig. 11). This is a good starting point to obtain diethyl 5-amino-1,3-oxazol-4-ylphosphonates (compounds 33 and 34). Availability of the latter ones opened an original route to structurally variable phosphonopeptides 35 and 36.<sup>49–53</sup>

A preliminary study on the addition of thiols and amines to double bonds of quite readily affordable phosphono dehydropolypeptide 37 (Fig. 12) indicated that this procedure could be a new mean to produce phosphono peptides containing non-typical side chains (compounds 38 and 39).<sup>54,55</sup>

The peptide 37 is also a good entry for compounds 40 by means of addition–elimination reaction upon action of bromine.<sup>55</sup> Upon reaction of 40 with thiophenol functionalized dehydropolypeptide 41 was obtained. However, the further studies are required to validate general utility of these reactions.

## C Peptides with non-hydrolysable analogues of phosphoamino acids

Post-translational modification of proteins by phosphorylation is an integral component of cellular physiology. Hundreds of thousands of sites of protein phosphorylation have been discovered but the understanding of the functions of the vast majority of these post-translational modifications is lacking. In

that respect transfer of a phosphoryl group to an amino acid side chain and formation of phosphoamino acid residues (phosphoserine, phosphothreonine, phosphotyrosine, phosphohistidine, phospholysine and/or phosphoarginine) cause that they quite often serve as molecular switches to control protein–protein interactions.<sup>56</sup> Physiological functions of these cell cycle proteins and their relevance for cancer stimulate the studies on their physiologic role and makes them the possible targets in cancer control. Therefore, it is not surprising that syntheses and biological evaluation of non-hydrolysable mimics of these amino acids and their peptides is an active field of research.<sup>57–59</sup>

It is worth to notice that hydrolysis of natural antibiotics rhizocticins and plumbemycins in bacterial cells provides an analogue of phosphothreonine (compound 42, Fig. 13),<sup>20</sup> a potent inhibitor of threonine synthase, which exerts antibacterial action inside the cell. Unfortunately, this analogue, as well as its peptides, has never been tested as possible cell cycle proteins regulator.

Phosphoproteins and phosphopeptides are considered as useful tools for elucidating biological functions of phosphorylations. However, the hydrolytic liability of phosphates in a cellular context limits their use in such studies. Thus, phosphatases inert mimetics, in which phosphate methylene or

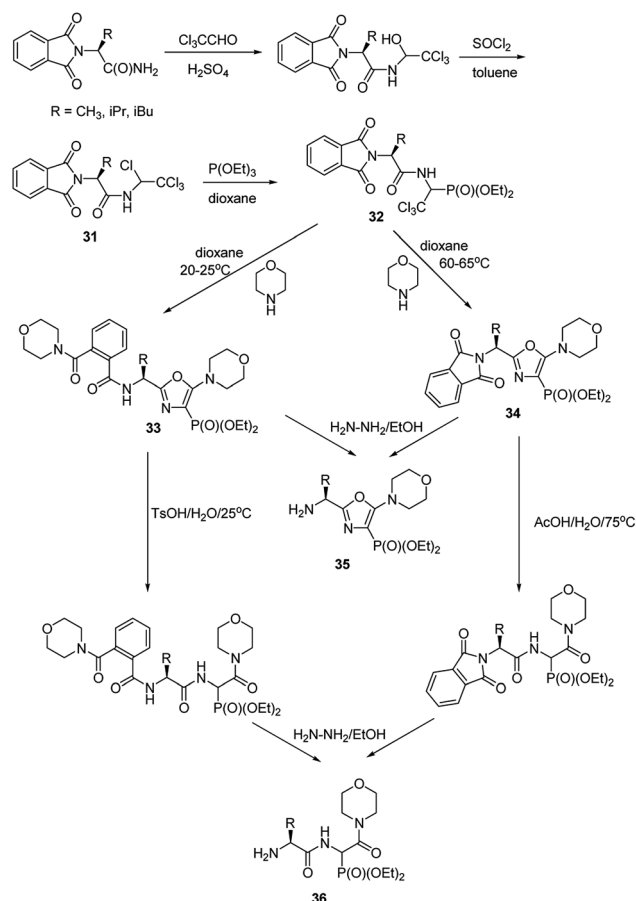


Fig. 11 Synthesis of phosphonopeptides via 5-amino-1,3-oxazol-4-ylphosphonates.

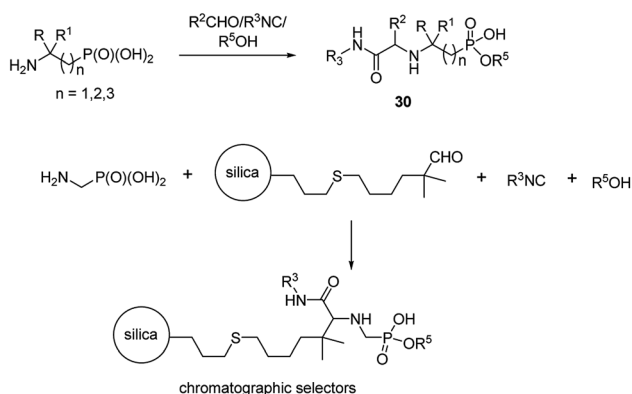


Fig. 10 Phosphonopeptides designed as phases for chromatography.

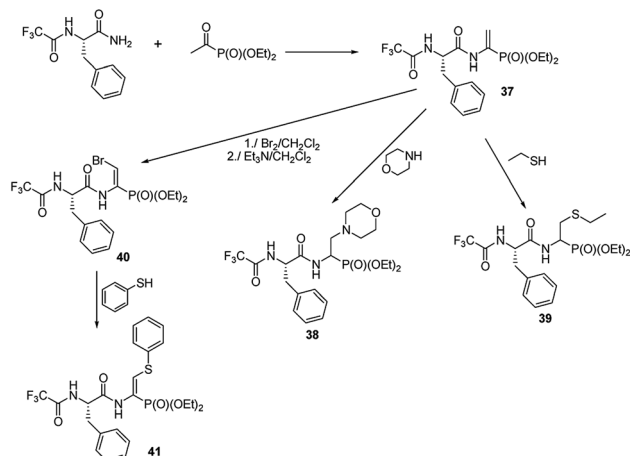


Fig. 12 Functionalization of phosphonodehydropeptide.

difluoromethylene groups replace oxygen, have been developed and used in many biological studies and therapeutic designs.<sup>59,60</sup> It is worth to note that difluoromethylene group is a better mimetic of the oxygen atom in phosphopeptides than methylene moiety in terms of basicity and charge distribution.<sup>61</sup>

The cell cycle regulatory serine/threonine-specific polo-like kinase is recognized as an oncogenic target. Peptides containing analogues, in which phosphate moiety of phosphothreonine is replaced by its non-hydrolysable analogue (compounds **43** and **44**, Fig. 13), appeared to be potent effectors of this cell cycle regulatory protein and may serve as a tool to study its role in cancerogenesis.<sup>62–65</sup> Masking the phosphonate moiety of peptide **43** by esterase labile pivaloyloxymethyl groups provided interesting prodrug.<sup>63</sup>

The major challenge in the studies on these mimetics is their synthesis. In most cases threonine and serine serve as substrates of choice in the elaborated procedures.<sup>66–69</sup> One of the most effective is shown in Fig. 14. Thus, methyl threoninate upon action of acetic acid anhydride was fully acylated and then converted to *Z* form of dehydroaminobutyrate (compound **45**). Reaction with *N*-bromosuccinimide (NBS) followed by treatment with strong base resulted in bromo derivative (compound **46**), which was coupled with diethyl difluoromethylenephosphonate providing *E*-prochiral substrate **47**. This compound was asymmetrically reduced using chiral

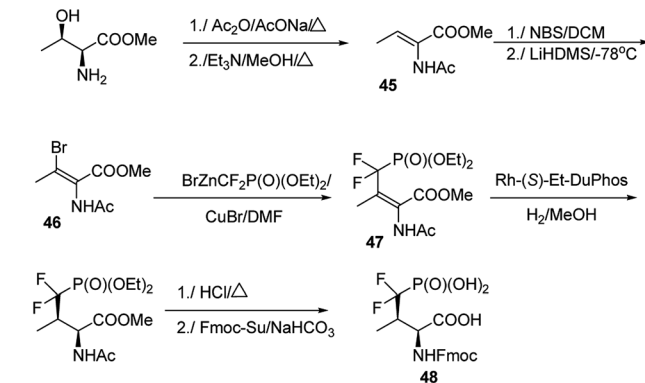


Fig. 14 Representative synthetic route to mimetic of phosphothreonine.

catalyst and finally converted into compound **48** suitable for introduction into peptide chain.<sup>64</sup> Preparation of the desired peptides was achieved easily by using classical procedures of both liquid- and solid-phase peptide syntheses.

The 14-3-3 proteins were the first phosphoserine/phosphothreonine binding proteins to be discovered. They are a family of highly conserved dimeric acidic proteins that are found in all eukaryotic organisms that regulate essentially every major cellular function. They integrate and control multiple signalling pathways and these interactions place them at the center of the signalling hub that governs critical processes in cancer, including apoptosis, cell cycle progression, autophagy, glucose metabolism, and cell motility. In human cells, this family of proteins consists of seven distinct but highly homologous 14-3-3 isoforms:  $\beta$ ,  $\epsilon$ ,  $\eta$ ,  $\gamma$ ,  $\sigma$ ,  $\tau$ ,  $\zeta$ . Therefore, peptides able to bind selectively these proteins have been selected and used for elucidation of physiologic roles of certain 14-3-3 proteins. Such molecules containing phosphatase-resistant phosphopeptide mimetics were synthesized and their affinities to isoforms:  $\zeta$  (peptides **49** and **50**, Fig. 15), and  $\beta$  (peptide **51**) had been evaluated.<sup>61,66–70</sup> They might be used as valuable *in vivo* tools for probing the role of phosphorylations.

Another possibility is genetically directed incorporation of phosphonic mimetics into proteins involved in cell regulation processes. Preliminary studies indicate that it is still challenging task.<sup>71,72</sup>

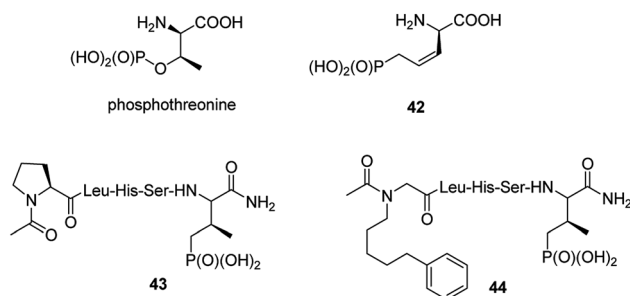


Fig. 13 Mimetics of phosphothreonine and its peptides.

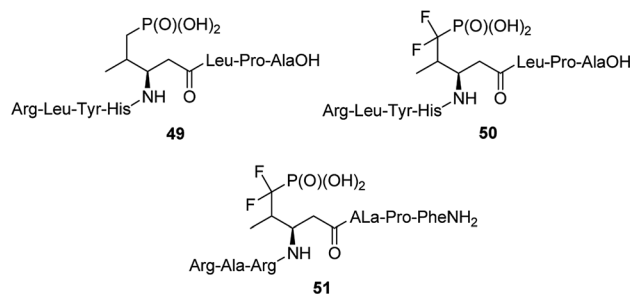


Fig. 15 Peptides containing non-hydrolysable mimetics of phosphotyrosine designed to bind 14-3-3-proteins.

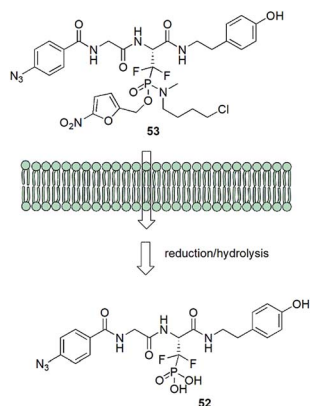


Fig. 16 Prodrug design for transportation of affinity probe through cell membranes.

Albeit being effective *in vitro* low-molecular weight inhibitors of phosphoserine/phosphothreonine binding proteins could not be used in cellular systems. It is mainly because the intracellular delivery of drugs that contain the necessary doubly charged phosphonate group is difficult because of the character of cell membrane. In order to get around this problem a very elegant system of transportation of affinity probe **52** was designed (Fig. 16). Ester moiety of the prodrug **53** is readily cleaved in the cell by enzymatic reduction followed by the hydrolysis of protonated phosphonamide and release of suitable inhibitor.<sup>62,69</sup>

Penta- to decapeptides containing non-hydrolysing analogue of pyrophosphoserine (see the representative structure **54** in Fig. 17) were obtained by solid-phase synthesis from bisphosphonate analogue **55**. They exerted enhanced stability in cell lysates and plasma, and therefore may serve as tools for studying, the marginally characterized to date, role of pyrophosphorylation.<sup>73</sup> In order to generate antibodies they were ligated with bovine serum albumin. The ligations were carried out: (i) by reaction of mimetic-hydrazide with N-terminal cysteine of chosen oligopeptide followed by radical desulfurization (formation of alanine); (ii) by reaction of N-terminal cysteine of the mimetic peptide with maleimide-activated bovine albumin.<sup>73</sup>

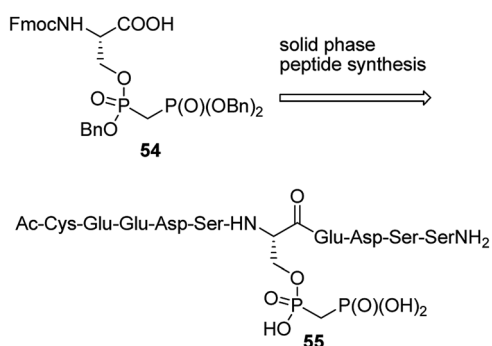


Fig. 17 Peptides containing mimetic of pyrophosphate-substituted serine.

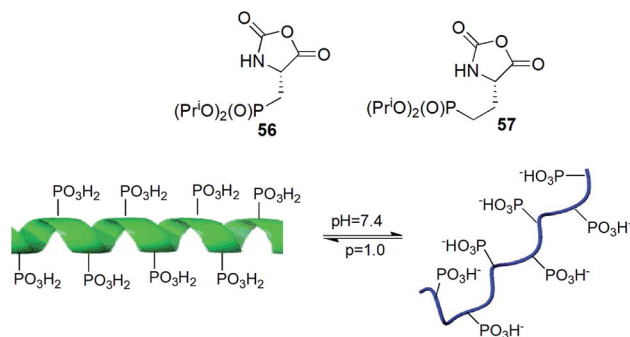


Fig. 18 pH-responsive polymers of phosphoserine mimetics.

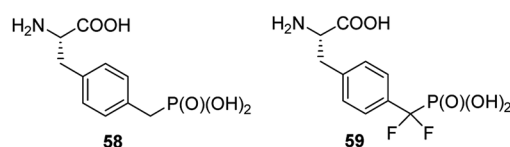


Fig. 19 Analogues of phosphotyrosine designed to study Src SH2 homology domains and tyrosine phosphatases.

*N*-Carboxyanhydrides of diisopropyl phosphonoalanine and homoalanine (compounds **56** and **57**) underwent metal-catalysed living polymerization yielding peptides of narrow chain length distribution and chain lengths to over 150 residues (Fig. 18). Copolymerization of compound **56** with appropriate *N*-carboxyanhydride of protected lysine provided block copolymers in high yields and low polydispersity indices. The obtained peptide exhibited pH-responsive conformational changes (Fig. 18).<sup>74</sup>

Non-hydrolysable phosphonic mimetics of phosphotyrosine were designed as molecular probes to study Src SH2 homology domains and tyrosine phosphatases. *p*-(Phosphonomethyl)-*L*-phenylalanine (compound **58**, Fig. 19) and its difluoro-derivative **59** are the most popular here.<sup>75,76</sup> The latter one better mimics electronic structure and acidity of mother counterpart. In last decade, however, the number of papers devoted to phosphonic analogues of phosphotyrosine decreased significantly and they relate rather to design of new effectors of specific proteins or their use in construction of diagnostic tools.

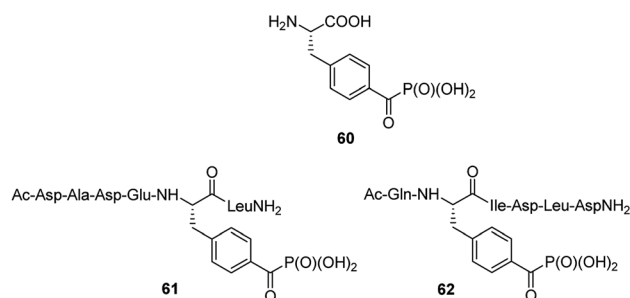


Fig. 20 Photoactive analogue of phosphotyrosine and its peptides.



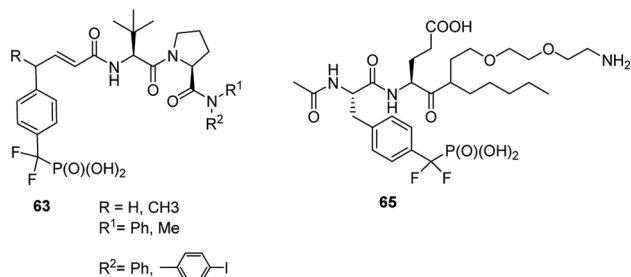


Fig. 21 Peptides containing non-hydrolysable analogue of phosphotyrosine.

Benzoylphosphonate analogue **60** (Fig. 20) appeared to act as photoactive, stable isoster of phosphotyrosine. It was introduced into two inhibitors (compounds **61** and **62**, Fig. 20), which are derived from phosphopeptide substrates of protein tyrosine phosphatase PTP1B. Photo-activation of these inhibitors results in the formation of triplet biradical intermediates of the benzoylphosphonate moiety, which deactivate PTP1B by an oxidative radical oxidation of active site Cys125.<sup>77</sup> These peptides if used in cell lysates or even living cells might possibly be useful tools in the time- and spatially resolved switching of PTP activities.

In response to cytokines and growth factors, STAT protein family members are phosphorylated by the receptor-associated kinases, and then form homo- or heterodimers that translocate to the cell nucleus where they act as transcription activators.

STAT6 plays a central role in exerting interleukins 4 and 13 (IL-4 and IL-13) mediated biological responses and thus modulate immunological response. Specific inhibitor of STAT6 containing compounds **63** (Fig. 21) were designed and synthesized.<sup>78</sup> They block recruitment to phosphotyrosine residues on IL-4 or IL-13 receptors and subsequent Tyr641 phosphorylation, which in turn results in inhibition of the expression of genes contributing to asthma.

Kv1.3 is one of 76 human potassium homotetrameric channels, which regulate membrane potential and calcium signaling in human T cells. It is considered as target for therapeutics

for multiple sclerosis. Recently a 35-residue disulfide-rich ShK peptide that blocks the voltage-gated potassium channels Kv1.3 and Kv1.1 has been isolated from sea anemone *Stichodactyla helianthus*. Modification of N-terminus of this peptide with *p*-(phosphonomethyl)phenylalanine (conjugate **64**, Fig. 22) resulted in highly selective picomolar blocker of this channel.<sup>79</sup>

Identification of particular SH2 domains is of importance since it may be used to shape the diagnosis. Consequently, the first step here is the enrichment of all (or at least majority) SH2 proteins present in biological material. Constructing an affinity resin containing general affinity probe could be a resolution of this problem. The preliminary studies with compound **65** (Fig. 21) have shown that this is possible since this compound was used for successful enrichment of 22 SH2 proteins from mixed cell lysates containing 50 of these proteins.<sup>80</sup>

Phosphorylation of histidine is recognized as being critical to signaling processes in bacteria, fungi and plants. Their role in eukaryotes is far less understood. In order to overcome the problems of low stability and isomerism observed for phosphohistidine (Fig. 24) their stable analogues have been synthesized and used as tools (compounds **66–71**, Fig. 23). Their use and utility has been recently reviewed.<sup>81–83</sup>

One of the drawbacks in studies on the role of histidine phosphorylation in mammals is a lack of specific antibodies to both isomers of phosphohistidine. Some studies on the use of phosphonate mimics of isomers of phosphohistidine have already been published. Such antibodies were raised by using oligopeptides containing these mimetics conjugated with bovine serum albumin (BSA)<sup>84,85</sup> and keyhole limpet hemocyanin (KLH).<sup>86,87</sup> The structures of the representative antigens obtained are shown in Fig. 24. The preliminary results of the cell studies are promising.

Phosphohistidine has two isomeric forms (1-phosphohistidine and 3-phosphohistidine, Fig. 25) and both of them exist in nature. Phosphoryltriarylalanine analogues (compounds **69** and **70**, Fig. 23) represent nonhydrolysable and non-isomerisable analogues of these isomers and both were used for construction of the above-mentioned antibodies (Fig. 24). They are readily available thanks to application of Huisgen dipolar cycloaddition (click reaction) of the commercially available ethynylphosphonate (compound **72**) and N-protected serine-derived azides (for example, Fmoc protected compound **73**). Regioselectivity of the reaction is strongly affected by applied catalyst and, to a lesser extent, by carboxy-protected group

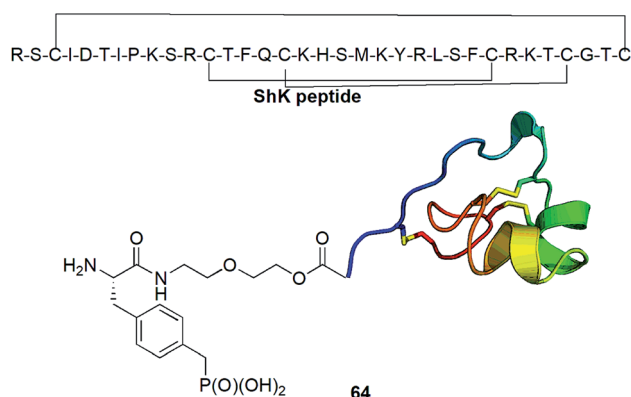


Fig. 22 Modified *Stichodactyla helianthus* peptide as potential agent against multiple sclerosis.

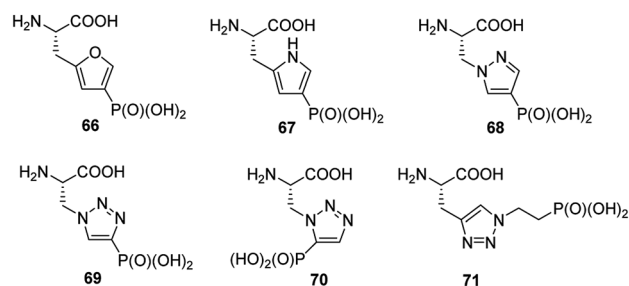


Fig. 23 Non-hydrolysable analogues of phosphohistidine.





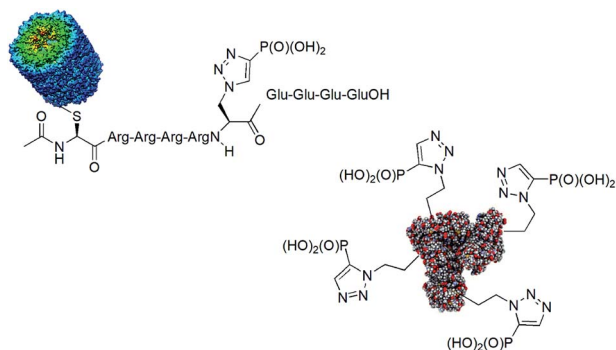


Fig. 24 Representative antigens containing mimetics of phosphohistidine.

(Fig. 25). Importantly, click reaction could be also applied using peptides containing diaminopropionic acid inside the peptide chain.<sup>81,88,89</sup>

Synthesis of phosphonopyrazol-4-yl alanine (compound **68**, Fig. 23) represents second type of general syntheses of phosphohistidine mimetics (Fig. 26). It starts from the ring opening of the Vederas  $\beta$ -lactone **74** with appropriate histamine analogue (in this case 4-iodopyrazole) yielding the acid **75**, which after protection is phosphorylated by means of Hirao cross-coupling reaction. The two-step deprotection of the resulting product (compound **76**) yields desired compound **68**.<sup>87</sup>

Homologue of phosphonic mimetic of phosphohistidine (compound **77**, Fig. 27) was found to be inactive. However, comparison of its structure with phosphotyrosine revealed that it might be an interesting mimetic of the latter. Indeed after inclusion into peptide chain (compound **78**, Fig. 27) it was found to bind SH2 domain of eukaryotic cell signalling protein Grb2. Since its synthesis is relatively simple it is considered as more accessible mimetic of phosphotyrosine than current alternatives.<sup>89</sup>

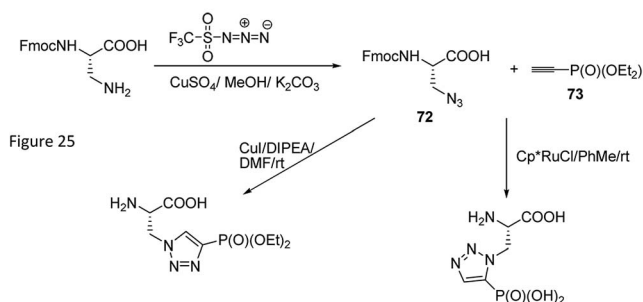
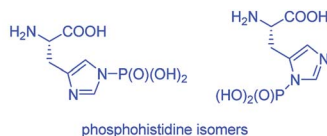


Fig. 25 Synthesis of analogues of phosphohistidine *via* click reaction.

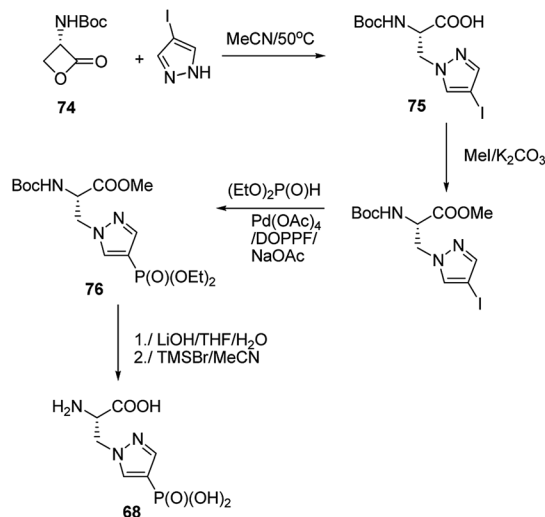


Fig. 26 Synthesis of phosphohistidine analogue **68**.

## D Peptides and proteins functionalized with phosphonic acid group

In this paragraph some dispersed data on synthesis of peptides and proteins containing phosphonic acid groups bound *via* different linkers is described. These entities were usually designed on purpose and were devoted to study selected physiologic processes.

CheY protein is involved in the transmission of sensory signals from the chemoreceptors to the bacterial flagellar motors because its binding to a switch component induces a change from counterclockwise to clockwise flagellar rotation and thus influence of bacterial chemotaxis. Better understanding of this process should lead to additional insights into bacterial pathogenicity. Phosphorylation of CheY promotes this association. However, biochemical studies of activated CheY-phosphate have been challenging due to the rapid hydrolysis of the aspartyl-phosphate *in vitro*. Therefore, mutated CheY, containing cysteine in place of active aspartyl acid, was alkylated with phosphonomethyltriflate providing non-hydrolysable analogue of phosphorylated CheY (conjugate **79**, Fig. 28).<sup>90</sup> It was found to bind effectively to the switch component in similar manner as CheY-phosphate.

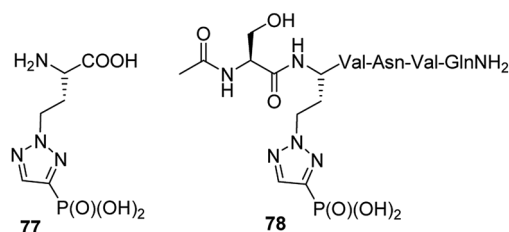


Fig. 27 Peptide containing analogue of phosphohistidine **76**, which acts as mimic of phosphotyrosine.

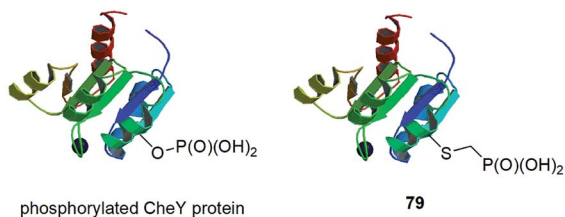


Fig. 28 CheY protein functionalized with phosphorus fragments.

Functionalization of peptides with bisphosphonic acid rises as a standard procedure for delivery of drugs to bone tissue. Thus, conjunction of potent inhibitor of proteasome with potent antiosteoporotic alendronate yielded two types of compounds: **80** and **81** (Fig. 29) designed as drugs against multiple myeloma,<sup>91</sup> whereas conjunction of doxorubicin with bisphosphonate using peptidyl linker provided compound **82**, which is a potential drug against bone metastases.<sup>92</sup>

Polymers that conduct protons are of crucial importance in clean energy research, since they are an ideal material for constructing hydrogel gels or mimetics of photosynthesis. A series of diblock copolymers **83** (Fig. 30) of poly-*N*-(2-ethyl)hexylglycine and poly-*N*-phosphonmethylglycine upon hydration (8 water molecules per phosphonate moiety) effectively conduct protons, which makes them particularly attractive for electrochemical applications.<sup>93</sup>

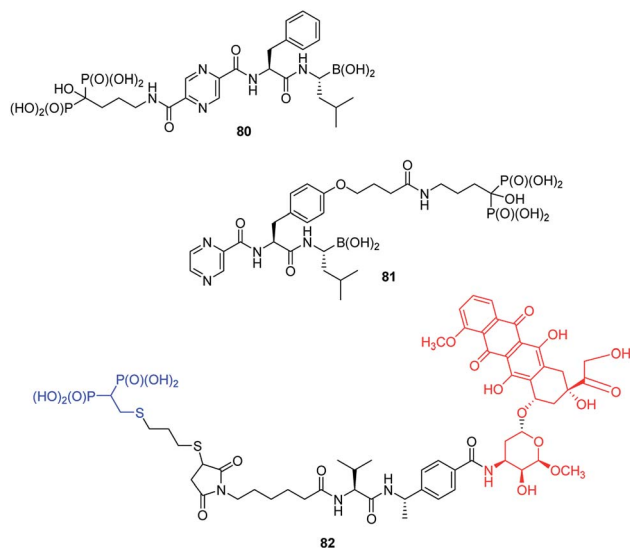


Fig. 29 Bisphosphonates as drug carriers.

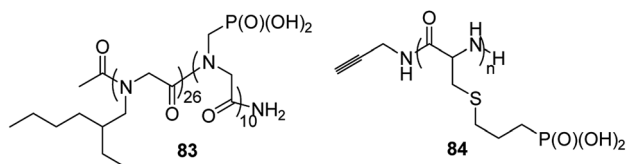


Fig. 30 Polymeric peptides of phosphonate holding amino acids.

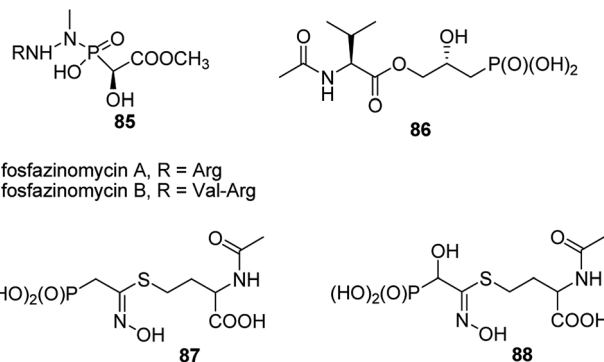


Fig. 31 Naturally occurring phosphonate peptidomimetics.

Phosphopolypeptides and their non-hydrolysable analogues are synthesized as mimetics of phosphorylated proteins that are involved in biomineralization. Such peptides (compounds **84**, Fig. 30)<sup>94</sup> were synthesized starting from *N*-propylphosphocysteine *N*-carboxyanhydride by simple polymerization upon action of primary amines.

A very specific cases worth to mention are: antifungal and antibacterial fosfazinomycins (compounds **85**, Fig. 31)<sup>95–97</sup> and valinophos (compound **86**).<sup>23</sup> Fosfazomycins are azapeptidyl analogues of phosphinopeptides and are rather representative of compounds with phosphonic group replacing peptide bond, whereas valinophos may be classified as phosphonodepsipeptide. Also recently isolated phosphonocystoximate and its hydroxyl-derivative (compounds **87** and **88**, Fig. 29)<sup>23</sup> could be considered as specific peptidomimetics.

## E Conclusions

Both groups of reviewed phosphono-peptides – those containing phosphonic acid moiety at C-terminus and those being non-hydrolysable mimetics of phosphonoamino acids, were discovered in 70s last century. Progress of the studies on these compounds has been quite slow, however, in last decade significant acceleration was noticed. In the case of antibacterial peptides it resulted from both, discovery of novel antibiotics by innovative and elegant genomic approach and from repurposing of old or abandon antibacterial agents, which is stimulated by fast growth of number of resistant bacterial strains. Effective methods for the synthesis of these compounds had been elaborated and optimized in last century. However, they require accessibility to esters of aminophosphonic acids, which usually are not readily available and have to be synthesized by independent procedures. Thus, modifications of simple phosphono-peptide side chains might be considered as useful alternative.

In the case of analogues of phosphoamino acids their possible functioning as molecular probes and generation of specific antibodies used as tools for more detailed study on the role and physiological consequences of phosphorylations at cellular level is a driving force of their synthesis. Thus, research described so far was concentrated on the elaboration of simple



and effective procedures for the synthesis of structurally variable mimetics.

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

The National Science Centre of Poland (NCN) grant number 2016/21/B/ST5/00115, supported this work.

## Notes and references

- 1 *Aminophosphonic and Aminophosphinic Acids: Chemistry and Biological Activity*, ed. V. Kukhar and H. Hudson, John Wiley & Sons, New York, 2000.
- 2 J. G. Allen, F. R. Atherton, M. J. Hall, C. H. Hassal, S. W. Holmes, R. W. Lambert, L. J. Nisbet and P. S. Ringrose, *Nature*, 1978, **272**, 56–58.
- 3 E. Bayer, K. H. Gugel, K. Hägele, H. Hagenmaier, S. Jessipow, W. A. König and H. Zähler, *Helv. Chim. Acta*, 1972, **55**, 224–239.
- 4 Y. Ogawa, T. Tsuruoka, S. Inouye and T. Niida, *Sci. Rep. Meiji Seika Kaisha*, 1973, **13**, 42–48.
- 5 M. Maślanka and A. Mucha, *Pharmaceuticals*, 2019, **12**, 86.
- 6 E. Burchacka and M. Sieńczyk, *Curr. Pharm. Des.*, 2018, **37**, 4445–4465.
- 7 R. Grzywa and M. Sieńczyk, *Curr. Pharm. Des.*, 2013, **19**, 1154–1178.
- 8 A. Mucha, P. Kafarski and Ł. Berlicki, *J. Med. Chem.*, 2011, **54**, 5955–5980.
- 9 A. Łupicka-Słowik, M. Psurski, R. Grzywa, M. Cuprych, J. Ciekot, W. Goldeman, E. Wojaczyńska, J. Wojaczyński, J. Oleksyszyn and M. Sieńczyk, *Invest. New Drugs*, 2020, **38**, DOI: 10.1007/s10637-020-00923-4.
- 10 S. McGowan, C. J. Porter, J. Lowther, C. M. Stack, S. J. Golding, T. S. Skinner-Adams, K. R. Trenholme, F. Teuscher, S. M. Donnelly, J. Grembecka, A. Mucha, P. Kafarski, R. Degori, A. M. Buckle, D. L. Gardiner, J. C. Whisstock and J. P. Dalton, *Proc. Natl. Acad. Sci.*, 2009, **106**, 2537–2542.
- 11 T. R. Burke, M. S. Smyth, M. Nomizu, A. Otaka and P. R. Roller, *J. Org. Chem.*, 1993, **58**, 1336–1340.
- 12 M. Talma and A. Mucha, *Biomolecules*, 2020, **10**, 659.
- 13 M. Talma, M. Maślanka and A. Mucha, *Bioorg. Med. Chem. Lett.*, 2019, **29**, 1031–1042.
- 14 M. M. Abdou, P. M. O'Neill, E. Amigues and M. Matziari, *Drug Discovery Today*, 2019, **24**, 916–929.
- 15 D. Georgiadis and V. Dive, *Top. Curr. Chem.*, 2015, **360**, 1–38.
- 16 A. Mucha, *Molecules*, 2012, **17**, 13530–13568.
- 17 F. R. Atherton, M. J. Hall, C. H. Hassal, R. W. Lambert, W. J. Lloyd and P. S. Ringrose, *Antimicrob. Agents Chemother.*, 1979, **15**, 696–705.
- 18 M. Leason, D. Cunliffe, D. Parkin, P. J. Lea and B. J. Mifflin, *Phytochemistry*, 1982, **21**, 855–857.
- 19 S. Omura, M. Murata, H. Hanaki, K. Hinotozawa, R. Oiwa and H. Tanaka, *J. Antibiot.*, 1984, **37**, 829–835.
- 20 M. Gahungu, A. Arguelles-Arias, P. Fickers, A. Zervosen, B. Joris, C. Damblon and A. Luxen, *Bioorg. Med. Chem.*, 2013, 4958–4967.
- 21 R. D. Johnson, R. M. Kastner, S. H. Larsen and E. E. Ose, *US Pat.* 4,482,488, 1984.
- 22 B. T. Circello, C. G. Miller, J.-H. Lee, W. A. van der Donk and W. W. Metcalf, *Antimicrob. Agents Chemother.*, 2011, **55**, 3357–3362.
- 23 K.-S. Ju, J. Gao, J. R. Doroighazi, K.-K. A. Wang, C. J. Thibodeaux, S. Li, E. Metzger, J. Fudala, J. Su, J. K. Zhang, J. Lee, J. P. Cioni, B. S. Evans, R. Hirota, D. P. Labeda, W. A. van der Donk and W. W. Metcalf, *Proc. Natl. Acad. Sci.*, 2015, **112**, 12175–12180.
- 24 M. Yamato, T. Koguchi, R. Okachi, K. Yamada, K. Nakayama, H. Kase, A. Karasawa and K. Shuto, *J. Antibiot.*, 1986, **39**, 44–52.
- 25 G. J. Kramer, A. Mohd, L. S. U. Schwager, G. Masuyer, K. R. Acharya, E. D. Sturrock and B. O. Bachmann, *ACS Med. Chem. Lett.*, 2014, **5**, 346–351.
- 26 A. Mucha and P. Kafarski, *Methods Mol. Biol.*, 2020, **2103**, 287–301.
- 27 N. Rabasso and N. Fadel, *Phosphorus, Sulfur Silicon Relat. Elem.*, 2011, **186**, 1811–1819.
- 28 J. M. Cervera-Villanueva Jr, J. L. Viveros-Ceballos, I. Lizanga-Elizalde and M. Ordóñez, *J. Pept. Sci.*, 2015, **22**, 70–75.
- 29 K. V. Turcheniuk, K. O. Poliashko, V. P. Kukhar, A. B. Rozhenko, V. A. Soloshonok and A. F. Sorochnikov, *Chem. Commun.*, 2012, **48**, 11519–11521.
- 30 M. Cortes-Clerget, O. Gager, M. Monteil, J.-L. Pirat, E. Migianu-Griffoni, J. Deschamp and M. Lecouvey, *Adv. Synth. Catal.*, 2016, **358**, 34–40.
- 31 M. Cortes-Clerget, J. Jover, J. Dussart, E. Kolodziej, M. Monteil, E. Migianu-Griffoni, O. Gager, J. Deschamp and M. Lecouvey, *Chem.-Eur. J.*, 2017, **23**, 6654–6662.
- 32 M. O. Clarke, X. Chen, A. Cho, W. E. Delaney IV, E. Doerffler, M. Fardis, M. Je, M. Mertzman, R. Pakdaman, H.-J. Pyun, T. Rowe, C. Y. Yang, X. C. Sheng and C. U. Kim, *Bioorg. Med. Chem. Lett.*, 2011, **21**, 3568–3572.
- 33 E. C. L. Marss, L. Varadi, A. F. Bedernjak, K. M. Day, M. Gray, A. L. Jones, S. P. Cummings, R. J. Anderson and J. D. Perry, *Molecules*, 2020, **25**, 1445.
- 34 K. T. Ng, J. D. Perry, E. C. L. Marss, S. Orenga, R. J. Anderson and M. Grey, *Molecules*, 2020, **25**, 1557.
- 35 D. J. Bougioukou, C. P. Ting, S. C. Peck, S. Mukherjee and W. A. van der Donk, *Org. Biomol. Chem.*, 2019, **17**, 822–829.
- 36 M. M. Jiménez-Andreu, F. J. Sayago and C. Cativiela, *Eur. J. Org. Chem.*, 2018, 3965–3973.
- 37 V. P. Kukhar and V. D. Romanenko, in *Amino Acids, Peptides and Proteins in Organic Chemistry: Modified Amino Acids, Organocatalysis and Enzymes*, ed. A. B. Hughes, Wiley-VCH Verlag GmbH & Co, Weinheim, 2010, Ch. 5, vol. 2.
- 38 R. A. Shastri, *Chem. Sci. Trans.*, 2019, **8**, 359–367.
- 39 B. Sun and J. Xu, *Synthesis*, 2015, **47**, 1479–1487.
- 40 B. Sun and J. Xu, *J. Pept. Sci.*, 2015, **21**, 615–619.



- 41 J. Xu, *Phosphorus, Sulfur Silicon Relat. Elem.*, 2019, **194**, 487–492.
- 42 F. Meng, F. He, X. Song, L. Zhang, W. Hu and G. Liu, *Amino Acids*, 2012, **43**, 423–429.
- 43 B. Sun, M. K. Sundaraneedi, H. M. Southam, R. K. Poole, I. F. Musgrave, F. R. Keene and J. G. Collins, *Dalton Trans.*, 2019, **48**, 14505–14515.
- 44 J. Liu, P. Liao, J. Hu, H. Zhu, Y. Wang, Y. Li, Y. Li and B. He, *Molecules*, 2017, **16**, 238.
- 45 P. Liao, S.-Q. Hu, H. Zhang, L.-B. Xu, J.-Z. Liu, B. He, S.-G. Liao and Y.-J. Li, *Bull. Korean Chem. Soc.*, 2018, **39**, 300–304.
- 46 M. Drąg, E. Wiczerzak, M. Pawelczak, Ł. Berlicki, Z. Grzonka and P. Kafarski, *Biochimie*, 2013, **95**, 1640–1649.
- 47 A. F. G. Gargano, S. Buchinger, M. Kohout, W. Lindner and M. Lämmerhoffer, *J. Org. Chem.*, 2013, **78**, 10077–10087.
- 48 A. F. G. Gargano, T. Leek, W. Lindner and M. Lämmerhoffer, *J. Chromatogr. A*, 2013, **1317**, 12–21.
- 49 K. M. Kondratyuk, E. I. Lukashuk, A. V. Golovchenko, E. B. Rusanov and V. S. Brovarets, *Russ. J. Gen. Chem.*, 2012, **28**, 643–651.
- 50 K. M. Kondratyuk, E. I. Lukashuk, A. V. Golovchenko, I. V. Komarov, V. S. Brovarets and V. P. Kukhar, *Tetrahedron*, 2013, **69**, 6251–6261.
- 51 O. I. Lukashuk, E. R. Abdurakhmanova, K. M. Kondratyuk, O. V. Golovchenko, K. V. Khokhlov, V. S. Brovarets and V. P. Kukhar, *RSC Adv.*, 2015, **5**, 11198–11206.
- 52 E. I. Lukashuk, E. R. Abdurakhmanova, K. M. Kondratyuk, A. V. Golovchenko and V. S. Brovarets, *Russ. J. Gen. Chem.*, 2015, **85**, 71–74.
- 53 E. R. Abdurakhmanova, O. I. Lukashuk, A. V. Golovchenko and V. S. Brovarets, *Russ. J. Gen. Chem.*, 2016, **86**, 1206.
- 54 B. Vanderhoydonck and C. V. Stevens, *Tetrahedron*, 2007, **63**, 7679–7689.
- 55 P. Lenartowicz, B. Dziuk, B. Zarychta, M. Makowski and P. Kafarski, *Phosphorus, Sulfur Silicon Relat. Elem.*, 2017, **192**, 706–712.
- 56 F. Ardito, M. Giuliani, D. Perrone, G. Troiano and L. Lo Muzio, *Int. J. Mol. Med.*, 2017, **40**, 271–280.
- 57 Z. Chen and P. A. Cole, *Curr. Opin. Chem. Biol.*, 2015, **28**, 115–122.
- 58 K. N. Chuh, A. R. Batt and M. R. Pratt, *Cell Chem. Biol.*, 2016, **23**, 86–107.
- 59 M. Arribat, F. Cavalier and E. Rémond, *RSC Adv.*, 2020, **10**, 6678–6724.
- 60 A. M. Marmelstein, J. Moreno and D. Fiedler, *Top. Curr. Chem.*, 2017, **375**, 20.
- 61 K. Panigrahi, D. L. Nelson and D. P. Berkovitz, *Chem. Biol.*, 2012, **19**, 666–667.
- 62 F. Liu, J.-E. Park, W.-J. Qian, D. Lim, A. Scharow, T. Berg, M. B. Yaffe, K. S. Lee and T. R. Burke, *ACS Chem. Biol.*, 2012, **7**, 805–810.
- 63 F. Liu, J.-E. Park, W.-J. Qian, D. Lim, A. Scharow, T. Berg, M. B. Yaffe, K. S. Lee and T. R. Burke, *ChemBioChem*, 2012, **13**, 1291–1296.
- 64 W.-J. Qian and T. R. Burke, *Amino Acids*, 2013, **45**, 1143–1148.
- 65 D. Hymel and T. R. Burke Jr, *ChemMedChem*, 2017, **12**, 202–206.
- 66 H.-Z. Duan, H.-X. Chen, Q. Yu, J. Hu, J.-M. Li and Y.-X. Chen, *Org. Biomol. Chem.*, 2019, **17**, 2099–2102.
- 67 H.-X. Chen, J. Kang, R. Chang, Y.-L. Zhang, H.-Z. Duan, Y.-M. Li and Y.-X. Chen, *Org. Lett.*, 2018, **20**, 3278–3281.
- 68 J. Kang, H.-X. Chen, S.-Q. Huang, Y.-L. Zhang, R. Chang, F.-Y. Li, Y.-M. Li and Y.-X. Chen, *Tetrahedron Lett.*, 2017, **58**, 2551–2553.
- 69 A. A. Arrendale, K. Kim, J. Y. Choi, W. Li, R. L. Geahlen and R. F. Borch, *ACS Chem. Biol.*, 2012, **19**, 764–771.
- 70 K. Ebisuno, M. Denda, K. Oguro, T. Inokuma, A. Shigenaga and A. Otaka, *Bioorg. Med. Chem.*, 2014, **22**, 2984–2991.
- 71 V. Beránek, C. D. Reinkemeier, M. S. Zhang, A. D. Liang, G. Kym and J. W. Chin, *Cell Chem. Biol.*, 2018, **25**, 1067–1074.
- 72 R. Klingberg, J. O. Jost, M. Schümann, K. A. Gelato, W. Fischle, E. Krause and D. Schwarzer, *ACS Chem. Biol.*, 2015, **10**, 138–145.
- 73 L. M. Yates and D. Fiedler, *ACS Chem. Biol.*, 2016, **11**, 1066–1073.
- 74 I. Yakovlew and T. J. Deming, *ACS Macro Lett.*, 2014, **3**, 378–381.
- 75 R. A. Cerulli and J. A. Kritzer, *Org. Biomol. Chem.*, 2020, **18**, 583–605.
- 76 P. Morlacchi, F. M. Robertson, J. Klostergaard and J. S. Mac Murray, *Future Med. Chem.*, 2014, **6**, 1909–1926.
- 77 S. Wagner, A. Schütz and J. Rademann, *Bioorg. Med. Chem.*, 2015, **23**, 2839–2847.
- 78 P. K. Mandal, P. Moriacchi, J. M. Knight, T. M. Link, G. R. Lee IV, R. Nurieva, D. Singh, A. Dhanik, L. Kavarak, D. B. Corry, J. E. Ladbury and J. S. McMurray, *J. Med. Chem.*, 2015, **58**, 8970–8984.
- 79 M. W. Pennington, S. C. Chang, S. Chauhan, R. Huq, R. B. Tajhya, S. Chhabra, R. S. Norton and C. Beeton, *Mar. Drugs*, 2015, **13**, 529–542.
- 80 M. Höfener, S. Heinzmler, B. Kuster and N. Sewald, *Proteome Sci.*, 2014, **12**, 41.
- 81 T. E. McAllister, M. G. Nix and M. E. Webb, *Chem. Commun.*, 2011, **47**, 1297–1299.
- 82 M. V. Makwana, R. Muimo and R. F. W. Jackson, *Lab. Invest.*, 2018, **98**, 291–303.
- 83 T. E. McAllister, J. J. Hollins and M. E. Webb, *Biochem. Soc. Trans.*, 2013, **41**, 1072–1077.
- 84 J.-M. Kee, R. C. Oslund, D. H. Perlman and T. W. Muir, *Nat. Chem. Biol.*, 2013, **9**, 416–421.
- 85 S. R. Fuhs, J. Meisenhelder, A. Aslanian, L. Ma, A. Zagorska, M. Stankova, A. Binnie, F. Al-Obeidi, J. Mauger, G. Lemke, J. R. Yates III and T. Hunter, *Cell*, 2015, **162**, 198–210.
- 86 J.-M. Kee, R. C. Oslund, A. D. Couvillon and T. W. Muir, *Org. Lett.*, 2015, **17**, 187–189.
- 87 M. Lilley, B. Mambwe, M. J. Thompson, R. F. W. Jackson and R. Muimo, *Chem. Commun.*, 2015, **51**, 7305–7308.
- 88 J.-M. Kee, B. Villani, L. R. Carpenter and T. W. Muir, *J. Am. Chem. Soc.*, 2010, **132**, 14327–14329.
- 89 T. E. McAllister, K. A. Horner and M. E. Webb, *ChemBioChem*, 2014, **15**, 1088–1091.





- 90 M. S. Beyersdorf, R. Sircar, D. B. Lookadoo, C. J. Bottone, M. J. Lynch, B. R. Crane and C. J. Halkides, *Protein Sci.*, 2017, **26**, 1547–1554.
- 91 J. K. Agyin, B. Santhamma and S. S. Roy, *Bioorg. Med. Chem. Lett.*, 2013, **23**, 6355–6458.
- 92 K. Hochdörffer, K. Abu Ajaj, C. Schäfer-Obodozie and F. Kratz, *J. Med. Chem.*, 2012, **55**, 7501–7512.
- 93 J. Sun, X. Jing, A. Siegmund, M. D. Conolly, K. H. Downing, N. P. Balsara and R. N. Zuckermann, *Macromolecules*, 2016, **49**, 3083–3090.
- 94 S. Das, M. Kar and S. S. Gupta, *Polym. Chem.*, 2013, **4**, 4087–4091.
- 95 S. Gunji, K. Arima and T. Beppu, *Agric. Biol. Chem.*, 1983, **41**, 2016–2069.
- 96 Z. Huang, K.-K. A. Wang and W. A. van der Donk, *Chem. Sci.*, 2016, **7**, 5219–5223.
- 97 J. Gao, K.-S. Ju, X. Yu, J. E. Velásques, S. Mukherjee, J. Lee, C. Zhao, B. S. Evans, J. R. Doroghazi, W. W. Metcalf and W. A. van der Donk, *Angew. Chem., Int. Ed.*, 2014, **53**, 1334–1337.

