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Evaluation of high-affinity phenyltetrahydroisoquinoline aldoximes, linked through *anti*-triazoles, as reactivators of phosphylated cholinesterases

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Abstract

Acetylcholinesterase (AChE) is a pivotal enzyme in neurotransmission. Its inhibition leads to cholinergic crises and could ultimately result in death. A related enzyme butyrylcholinesterase (BChE) may act in the CNS as a co-regulator in terminating nerve impulses and is a natural plasma scavenger upon exposure to organophosphate (OP) nerve agents that irreversibly inhibit both enzymes. With the aim of improving reactivation of cholinesterases phosphylated by nerve agents sarin, VX, cyclosarin, and tabun, ten phenyltetrahydroisoquinoline (PIQ) aldoximes were synthesized by Huisgen 1,3 dipolar cycloaddition between alkyne- and azide-building blocks. The PIQ moiety may serve as a peripheral site anchor positioning the aldoxime moiety at the AChE active site. In terms of evaluated dissociation inhibition constants, the aldoximes could be characterized as high-affinity ligands. Nevertheless, high binding affinity of these oximes to AChE or its phosphylated conjugates did not assure rapid and selective AChE reactivation. Rather, potential reactivators of phosphylated BChE, with its enlarged acyl pocket, were identified, especially in case of cyclosarin, where the reactivation rates of the lead reactivator was 100- and 6-times that of 2-PAM and HI-6, respectively. Nevertheless, the return of the enzyme activity was affected by the nerve agent conjugated to catalytic serine, which highlights the lack of the universality of reactivators with respect to both the target enzyme and OP structure.

Keywords

organophosphate; antidote; nerve agent; acetylcholinesterase; butyrylcholinesterase; pralidoxime; docking; peripheral anionic site

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1. Introduction

In the past decades, we have witnessed numerous cases of abuse of organophosphorus compounds (OPs) with nerve agents against civilians in war (Dolgin, 2013) or as means of terrorist attacks (Nakagawa et al., 2018; Vale et al., 2018) resulting in severe outcomes due to the irreversible inhibition of the pivotal enzyme acetylcholinesterase (AChE) and the related enzyme butyrylcholinesterase (BChE). The role of BChE is not essential, but it acts as a natural scavenger from tissue AChE inhibition by OPs and may serve as a CNS co-regulator of cholinergic neurotransmission (Raveh et al., 1993; Mesulam et al., 2002; Masson and Nachon, 2017). Currently used therapy in the event of OP poisoning, consisting of an oxime reactivator (2-PAM, obidoxime, HI-6), atropine and an anticonvulsant remains limiting due to variability in the oxime efficacy regarding different conjugated OPs (Gray, 1984; Dawson, 1994; Antonijevi and Stojiljkovi 2007; Worek et al., 2016). Over the years, numerous approaches were employed in order to improve the reactivation efficacy which is primarily attributed to the nucleophilic displacement rate of OPs and apparent affinity of the reactivating oxime for the phosphorylated cholinesterases (ChE). Therefore, efficacy varies with the bound OP and the structure of the oxime (Millard et al., 1999; Kovarik et al., 2004, 2006, 2019a; Ekström et al., 2006a, 2006b; Wille et al., 2010; Wong et al., 2000; Worek et al., 2002). Concepts for the development of new antidotes include miscellaneous ligands that form productive interactions with the ChE active centers and increases in enzyme binding affinity. Furthermore, a positive effect on reactivation efficiency was shown upon designing oximes that lack a permanent charge (Kalisiak et al., 2011; Sit et al., 2011, 2018; Radi et al. 2012; Kovarik et al., 2013; Rosenberg et al, 2017a, 2017b; Zorbaz et al., 2018a) or pro-drugs and amphipathic reactivators, like pro-2-PAM, that improve the delivery of the oxime to the central nervous system (Demar et al., 2010; Chambers et al., 2016).

The active site of ChE may be divided into two sub-sites: the catalytic site (CAS) at the base of the gorge, composed of a catalytic triad, oxyanion hole, acyl binding pocket, and choline binding site; and the peripheral anionic site (PAS) at the gorge rim (Taylor and Radi, 1994). The difference in amino acid residue composition of AChE and BChE active sites enables BChE to hydrolyze or bind larger substrates and ligands than AChE (Radi et al., 1993; Taylor et al., 1995; Saxena et al., 1999; Kovarik, 1999; Kovarik et al., 2003; Bosak et al., 2008). Encouraged by the hypothesis that the increase of the binding affinity of the ChE would enhance the reactivation efficacy, dual-binding mode reactivators containing ligand(s) that form interactions with both CAS and PAS (Ekström et al., 2009) or triple-binding mode compounds (Marakovi et al., 2016) were designed and tested. Researchers also found design inspiration in agents used in the treatment of Alzheimer's disease, like tacrine or donepezil, which interact with aromatic residues of AChE (Kryger et al., 1999; McHardy et al., 2014; Renou et al., 2016). An improved reactivation efficacy was shown for piperidine derivatives, tetrahydroacridine and tryptoline moieties containing reactivators, etc. (De Koning et al., 2011; Kliachyna et al., 2014; McHardy et al., 2014; Renou et al., 2014; Zorbaz et al., 2018a). Moreover, oximes comprising tetrahydroisoquinoline (TIQ) and phenyltetrahydroisoquinoline (PIQ) groups were presented as equal or more potent reactivators of VX- and tabun-inhibited AChE than standard oximes (Mercey et al., 2011, 2012; Wei et al., 2014, 2017; Zorbaz et al., 2018a). Further, Krasinski et al. (2005) reported

micromolar to femtomolar affinity of the PIQ-comprising AChE inhibitors synthesized by *in situ* click chemistry. This and the fact that the triazole moiety was observed earlier to contribute synergistically to the total energy of binding of high-affinity AChE inhibitors (Manetsch et al., 2004) encouraged us to design triazole aldoximes linked to the PIQ moiety. In our recent study with triazole containing oximes, we reported a successful reactivation of tabun and other nerve agents inhibited AChE (Kovarik et al., 2019a, 2019b).

Herein, we report the synthesis of ten potential reactivators containing PIQ moiety using Huisgen 1,3 dipolar cycloaddition forming *anti*-triazoles. The family of reactivating oximes was tested as individual reactivators of human AChE and BChE inhibited by sarin, VX, cyclosarin, and tabun. We also estimated the AChE and BChE binding affinity for PIQ aldoximes in terms of the oxime-enzyme dissociation constants (K_i). Computational docking was used to visualize important interactions with the native or OP-conjugated enzyme and for the overall interpretation of reactivation kinetics.

2. Materials and methods

2.1. Chemistry

Reagents and solvents were purchased from Aldrich (St. Louis, MO, USA), Acros (Morris Plains, New Jersey, USA) or GFS (Columbus, Ohio, USA) and used without further purification. Alkynes and azides were prepared according to procedures from the literature (Krasinski et al., 2005; Kovarik et al., 2019a). ^1H and ^{13}C NMR spectra were recorded on a Varian INOVA-400 spectrometer in CDCl_3 (7.26 ppm for ^1H , 77.00 ppm ^{13}C), d_6 -DMSO (2.50 ppm for ^1H , 39.43 ppm for ^{13}C) as standards.

Electrospray ionization mass spectrometry (ESI-MS) gave the following results: **1a**: Solid (343 mg, 89%). ESI MS $[\text{M}]^+$ for $\text{C}_{32}\text{H}_{39}\text{N}_6\text{O}_3$, calcd 555.3, found 555.4. **1b**: Solid (363 mg, 92%). ESI MS $[\text{M}]^+$ for $\text{C}_{33}\text{H}_{41}\text{N}_6\text{O}_3$, calcd 569.3, found 569.5. **1c**: Solid (371 mg, 92%). ESI MS $[\text{M}]^+$ for $\text{C}_{34}\text{H}_{43}\text{N}_6\text{O}_3$, calcd 583.3, found 583.5. **1d**: Solid (370 mg, 96%). ESI MS $[\text{M}]^+$ for $\text{C}_{32}\text{H}_{39}\text{N}_6\text{O}_3$, calcd 555.3, found 555.5. **1e**: Solid (380 mg, 96%). ESI MS $[\text{M}]^+$ for $\text{C}_{33}\text{H}_{41}\text{N}_6\text{O}_3$, calcd 569.3, found 569.4. **2a**: Solid (380 mg, 96%). ESI MS $[\text{M}]^+$ for $\text{C}_{33}\text{H}_{41}\text{N}_6\text{O}_3$, calcd 569.3, found 569.4. **2b**: Solid (391 mg, 97%). ESI MS $[\text{M}]^+$ for $\text{C}_{34}\text{H}_{43}\text{N}_6\text{O}_3$, calcd 583.3, found 583.5. **2c**: Solid (406 mg, 99%). ESI MS $[\text{M}]^+$ for $\text{C}_{35}\text{H}_{45}\text{N}_6\text{O}_3$, calcd 597.3, found 597.5. **2d**: Solid (390 mg, 99%). ESI MS $[\text{M}]^+$ for $\text{C}_{33}\text{H}_{41}\text{N}_6\text{O}_3$, calcd 569.3, found 569.5. **2e**: Solid (398 mg, 99%). ESI MS $[\text{M}]^+$ for $\text{C}_{33}\text{H}_{41}\text{N}_6\text{O}_3$, calcd 569.3, found 569.5.

2.2. Cholinesterase activity measurements

Recombinant human AChE, wild type, was prepared as described earlier (Cochran et al., 2011). Purified oligomeric BChE isolated from human plasma was kindly donated by Dr. David Lenz and the late Dr. Douglas Cerasoli, USAMRICD, Edgewood, MD.

For reactivation experiments, the enzyme was incubated with a ten-fold molar excess of nerve agents (NC Laboratory, Spiez, Switzerland) for an hour, fractionated on a Sephadex G-50 spin column (Roche Diagnostic GmbH, Mannheim, Germany), and then incubated with an oxime. Oxime stock solutions (100 or 10 mM) were made in DMSO and further

dilutions were prepared in water. At specified time intervals, an aliquot was 100-fold diluted and upon addition of ATCh residual enzyme activity was measured. The same procedure was applied to the uninhibited enzyme and control activity was measured in the presence of an oxime at concentrations used for reactivation. Reactivation screening was done at a given oxime concentration (10 μM for AChE and 100 μM for BChE) and was monitored up to 24 h. No spontaneous reactivation of phosphorylated cholinesterases was detected and enzyme was stable in 24 h. Detailed reactivation kinetics using a wider oxime concentration range of leads enabled the determination of reactivation constants: k_2 (maximal first-order reactivation rate constant), K_{ox} (apparent phosphorylated enzyme-oxime dissociation constant) and k_r (overall second-order reactivation rate constant). k_2 and K_{ox} were evaluated from the plot k_{obs} vs oxime concentration where (Kovarik et al., 2004; Ma ek Hrvat et al., 2018). k_r was the ratio of k_2 and K_{ox} .

Reversible inhibition of AChE and BChE was measured in the presence of aldoxime (0.004 - 20 μM). The enzyme-oxime dissociation constant (K_i) was evaluated from the Hunter-Downs equation and procedures described previously (Kovarik et al. 2006; Katalini et al., 2017).

2.3. Computational molecular docking

A three-dimensional structure of mouse AChE (PDB code: 5EHN; Bourne et al., 2016) and human BChE (PDB code: 3DJY; Carletti et al., 2008) were used for docking. Aldoxime structures were modeled and minimized using the MMFF94 force field implemented in ChemBio3D Ultra 12.0 (PerkinElmer, Inc., Waltham, MA). Discovery Studio 2017 R2 with CDOCKER docking protocol using a CHARMM force field (BioVia Discovery Studio 2017R2 software, San Diego, USA) generated for each aldoxime 20 poses in the active site gorge of AChE or BChE (Marakovi et al., 2016). Poses were scored and ranked according to the calculated CDOCKER energy from interactions between oxime and active site residues (Šinko, 2019).

3. Results

3.1. Synthesis

Aldoximes were synthesized using copper(I)-catalyzed azide-alkyne building block Huisgen's [2+3] cycloaddition (CuAAC) to form a linking 1,4-triazole as shown on Scheme 1:

Alkynes and azides used for the synthesis of these 10 compounds (Figure 1) were prepared as described previously (Krasinsky et al., 2005; Kovarik et al., 2019a). The general method for the synthesis of 1,4-triazole-oximes is described as follows. The alkyne (0.55 mmol) and the azide (0.55 mmol) were dissolved in the mixture of t-BuOH:H₂O, 2:1 (3.3 mL), then sodium ascorbate (21.8 mg, 0.11 mmol, 20% mol) and CuSO₄ (4.39 mg, 0.027 mmol, 5% mol) were added. The reaction was carried out for overnight at room temperature until LC/MS analysis showed no starting materials. Subsequently, the mixture was diluted with H₂O (10 mL) and a solution of NaPF₆ (185 mg, 1.1 mmol, 2 equiv.) in H₂O (5 mL) was

added. The mixture was cooled down to 0 °C and the hexafluorophosphate salt was separated, washed with H₂O and dried on high-vacuum.

3.2. Reversible inhibition of cholinesterases by aldoximes

The evaluated enzyme-oxime dissociation constants (K_i ; Table 1) in nanomolar (for AChE) and sub-nanomolar to micromolar range (for BChE) classify the PIQ aldoximes as high-affinity binding inhibitors. Although both enzymes had the highest affinity ($1/K_i$) for aldoxime **2a**, the affinity for AChE was about 12-times higher than that for BChE. Generally, most of the PIQ compounds could be considered as selective AChE inhibitors, since their inhibition of BChE was 5- (**1a**) to 955-fold (**1b**) less potent than for AChE. From a comparison of inhibition potencies and aldoxime structure no straightforward generalizations appear. The most potent AChE inhibitor (**2a**) differs from the less potent (**1d**) in length of linker having a 6- vs a 5-methylenes between the triazole ring and PIQ moiety, and *ortho*- vs *para*-positioned oxime group, respectively. Yet, aldoxime **1a**, an *ortho*- analog of **1d**, is among of the top three most potent AChE inhibitors. In the case of BChE, the K_i determined for the most potent BChE inhibitors (**2a** and **1a**) and their analogs with a *para*-positioned oxime group (**2d** and **2a**) differed 2- and 60-times, respectively. Interestingly, **1d** and **2c** had an almost identical K_i determined for BChE, even though they are structurally very different (shortest vs longest triazole containing linker and *para*- vs *ortho*-positioned oxime group, respectively). A similar situation as was found for oximes **1b** and **1c**.

3.3. Oxime-assisted reactivation of cholinesterases inhibited by nerve agents

Ten phenyltetrahydroisoquinoline (PIQ) aldoximes (Figure 1) were screened for reactivation of sarin-, cyclosarin-, VX- and tabun-inhibited human AChE and BChE at a given oxime concentration, 10 μ M for AChE and 100 μ M for BChE, and the results were sorted in terms of the observed first-order reactivation rate (k_{obs}) (Figure 2). If the maximal percentage of reactivation ($React_{max}$) reached in 24 h was less than 30%, k_{obs} was not calculated. In case of AChE reactivation PIQ aldoximes were not more efficient than HI-6, although nine, five and two aldoximes reactivated more than 50% in case of cyclosarin, sarin, and VX inhibition, respectively. Tabun-inhibited AChE or BChE was resistant to reactivation by any of tested aldoximes (results not shown). Phosphylated BChE conjugates were more prone to be reactivated with PIQ aldoximes. All PIQ aldoximes reactivated more than 60% of cyclosarin- and sarin-inhibited BChE, and 40 - 50% of BChE activity in the case of VX inhibition (Figure 2). Moreover, several PIQ aldoximes were more potent reactivators of phosphylated BChE than HI-6 or 2-PAM, and for these detailed kinetic parameters were determined.

Kinetic studies over a range of oxime concentrations were completed for the selected reactivators of cyclosarin- and sarin-inhibited BChE (Figure 3), and results were sorted in terms of k_r as an overall measure of efficiency (Table 2). The cyclosarin-inhibited BChE activity was restored with selected aldoximes, **1d** and **2d**, up to 80% within a short time, 10 or 30 min, respectively. Both aldoximes have similar binding affinities ($1/K_{ox}$) but the rate of dephosphylation (k_2) was two-fold lower for **2d**. It seems that the length of the triazole-containing linker affected the reactivation efficacy, because aldoximes **1d** and **2d** differ only

by a five versus six methylene chain linked between the triazole ring and PIQ moiety, respectively (Figure 1). A 6- or a 100-fold higher k_r with aldoxime **1d** than with HI-6 or 2-PAM, respectively, seems to be due to a low binding affinity of cyclosarin-conjugated BChE for standard oximes (Table 2).

In case of sarin, four selected aldoximes reactivated up to 90% of sarin-inhibited BChE within 3 hours. Although high percentage of reactivation was achieved with both standard oximes (Table 2), and 2-PAM had the highest potency for the nucleophilic displacement of sarin moiety, its overall reactivation potency was lower than that of tested oximes due to its low apparent binding affinity (high K_{ox}). It is interesting to note that if one compares spacer length, shorter linker analogues (**1b** and **1e**) showed faster reactivation rate but lower affinity than their counterparts (**2b** and **2e**).

3.4. Molecular modelling of aldoxime complexes with AChE and phosphonylated BChE

The model of the complex between aldoxime **2a**, the most potent AChE inhibitor, and mouse AChE is shown in Figure 4. It seems that the high affinity of native AChE for **2a** is due to the multiple interactions of the aldoxime. The pyridinium ring of **2a** was stabilized by Trp86 from the choline binding site and by His447, a residue of the catalytic triad, while the oxime group was close to the catalytic Ser203. H-bonding with Gly121 from the oxyanion hole is also a possibility. The PIQ of **2a**, as expected, was located in the PAS, forming π - π interactions with Trp286, and H-bonds with Ser293 and Tyr341.

Aldoxime **1d**, a potent reactivator of cyclosarin-inhibited BChE, was modelled inside the model of BChE-cyclosarin conjugate using flexible docking allowing Leu286, Phe329, and Tyr332 to rotate freely. The minimised structure of the complex, presented in Figure 5 showed a favourable position of the oxime group for a nucleophilic attack with a distance of 4.6 Å from the phosphorus atom. The PIQ domain of **1d** was stabilized outside the choline-binding site of BChE via multiple H-bonds involving Trp82, Asn83, Thr120, Pro285, and Asn289. The triazole ring of **1d** was positioned close to Ala277. Due to a poorly defined PAS in BChE, lacking aromatic side chains, the conformation of **1d** was so oriented to be stabilized by an intramolecular cation- π and π - π interaction between the pyridinium ring and PIQ moiety of **1d**.

4. Discussion

This study describes the design, synthesis and *in vitro* evaluation of the interactions of ten new aldoximes and, their potential as reactivators of cholinesterases inhibited by the nerve agents: sarin, cyclosarin, VX, and tabun. Oximes were synthesized using Huisgen 1,3 dipolar cycloaddition forming *anti*-triazoles. Alkyne building blocks comprising the PAS-binding ligand PIQ should anchor the compound in a way that the azide building block with a pyridinium ring and oxime group is directed towards phosphate or phosphonate at the catalytic serine. The triazole ring formed by this cycloaddition should contribute to binding interactions within the active site gorge and additionally enhance the affinity of the enzyme for the compounds. This hypothesis was supported by the dissociation inhibition constants of AChE which were in nanomolar range (Table 1) classifying the tested aldoximes as its high-affinity ligands. Moreover, the molecular modeling study with aldoxime **2a** confirmed

the stabilization of its PIQ moiety in the PAS of AChE. Nevertheless, although we succeeded in improving the binding affinity of the native AChE for tested compounds, the efficacy of PIQ aldoximes as phosphorylated AChE reactivators was not superior to HI-6. This may indicate that oximes are stabilized in an unfavorable position for efficient nucleophilic attack. It seems that triazole in linker affect the reactivation, since other PIQ aldoximes have been previously reported as equally or more potent reactivators than the standard oximes for AChE inhibited by various OPs (Mercey et al., 2011, 2012; Zorbaz et al., 2018a). This may indicate that the triazole linkage as a strong dipole confers a less favorable oxime orientation for nucleophilic attack, despite the high affinity of these compounds for unconjugated AChE.

Conformational changes and steric hindrance in the gorge induced by the OP conjugation (Eto, 1976; Millard et al., 1999; Ekström et al., 2006a, 2006b; Katalini et al., 2018) consequently affect the possibility of the oxime to adopt the most favorable position for dephosphorylation of catalytic serine. This was particularly noted in the case of tabun inhibition where there was no reactivation of either AChE or BChE with the tested aldoximes. Yet, in our recent paper (Kovarik et al., 2019a) we reported a potent reactivator (oxime **3A**) of a phosphoramidated (tabun-AChE) conjugate, also synthesized by the Huisgen cycloaddition, but with a much shorter triazole containing linker between two pyridinium rings, when compared to PIQ aldoximes. On the other hand, although the active site of BChE allows more conformational freedom and flexibility of tested PIQ aldoximes than the active site of AChE, potential reactivators of cyclosarin- and sarin-, but not of VX- and tabun-inhibited BChE were identified emphasizing the complexity of reactivation as a consequence of the steric constraints of conjugated phosphorus moiety. It is interesting to note, that aldoxime **1d** is more efficient reactivator of the cyclosarin-BChE conjugate than a PIQ containing aldoxime reported by Zorbaz et al. (2018a). This is probably due to its higher binding affinity and stabilization of the triazole ring in the cyclosarin-BChE conjugate, as shown by molecular modelling (Figure 5).

The determined dissociation inhibition constants of BChE were in micromolar or submicromolar range, and if compared to the nanomolar constants with these compounds for AChE, the tested aldoximes are preferably complexed by AChE. Furthermore, detailed kinetics of the reactivation of cyclosarin- and sarin-inhibited BChE showed that the affinity of the phosphorylated BChE conjugate for top reactivators decreased up to 400 times, when compared to native BChE (Table 1, Table 2). This is probably due to the conformational changes in the active site and steric hindrance with the binding of the OP which affects oxime embedding and orientation. Therefore, achieving a moderate apparent affinity optimizes reactivation and, excessively high affinity does not necessarily enhance reactivation rates. Multiple interactions of oxime with active site residues could stabilize the oxime in unproductive conformation, with the oximate directed away from the phosphorus conjugated to catalytic serine, therein not fulfilling the alignment for efficient dephosphorylation. Alternatively, a spatially constrained active center gorge may restrict the angle of attack and the departure of the conjugated organophosphate from the gorge. Finally, the high affinities of these oxime reactivators for the native enzyme are likely to inhibit the residual active enzyme in an OP exposure interval. Accordingly, achieving maximal

reactivation *in vivo* and minimal toxicity from the reactivator itself is best be achieved with a maximal reactivation rate and minimizing the affinity for the residual free enzyme.

To conclude, the efficacy of reactivator reveals an interplay between affinity for the residual free and conjugated enzymes, the optimal length of the linker (compound), position and orientation of the oximate with respect to the conjugated phosphorus. Moreover, one must consider the structure of the phosphorylating agents since they also differentially restrict access to the gorge.

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Highlights

- phenyltetrahydroisoquinoline moiety acts as peripheral anionic site ligand
- potential reactivators of phosphorylated BChE were identified
- high binding affinity of AChE for oximes does not ensure efficient reactivation *in situ* oxime efficacy depends on the organophosphate conjugated to the cholinesterase

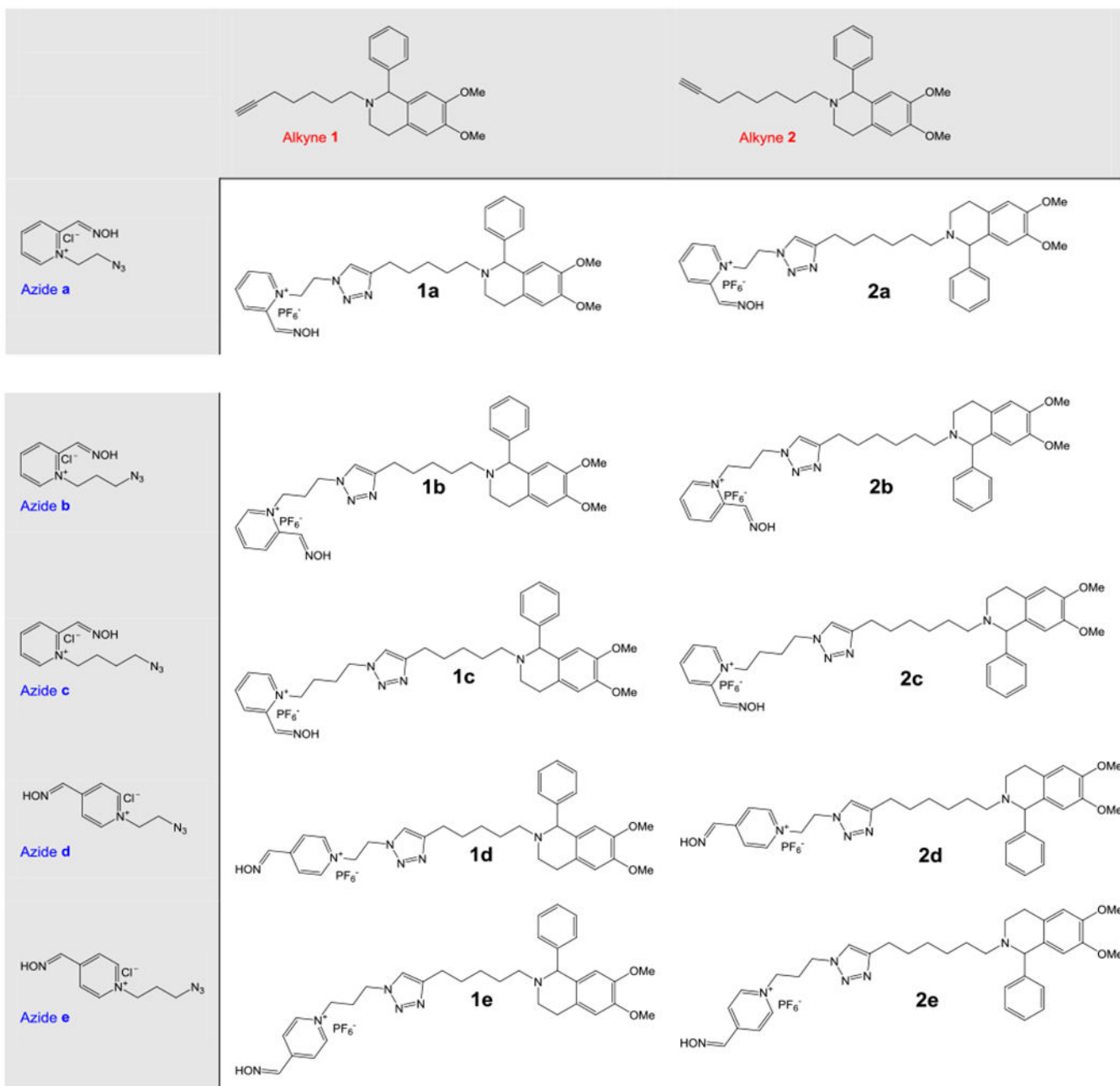


Figure 1. Chemical structures of alkyne (**1-2**) and azide (**a-e**) building blocks in copper(I)-catalyzed azide-alkyne building block [2+3] cycloaddition utilized for the synthesis of phenyltetrahydroisoquinoline aldoximes (**1a-1e** and **2a-2e**).

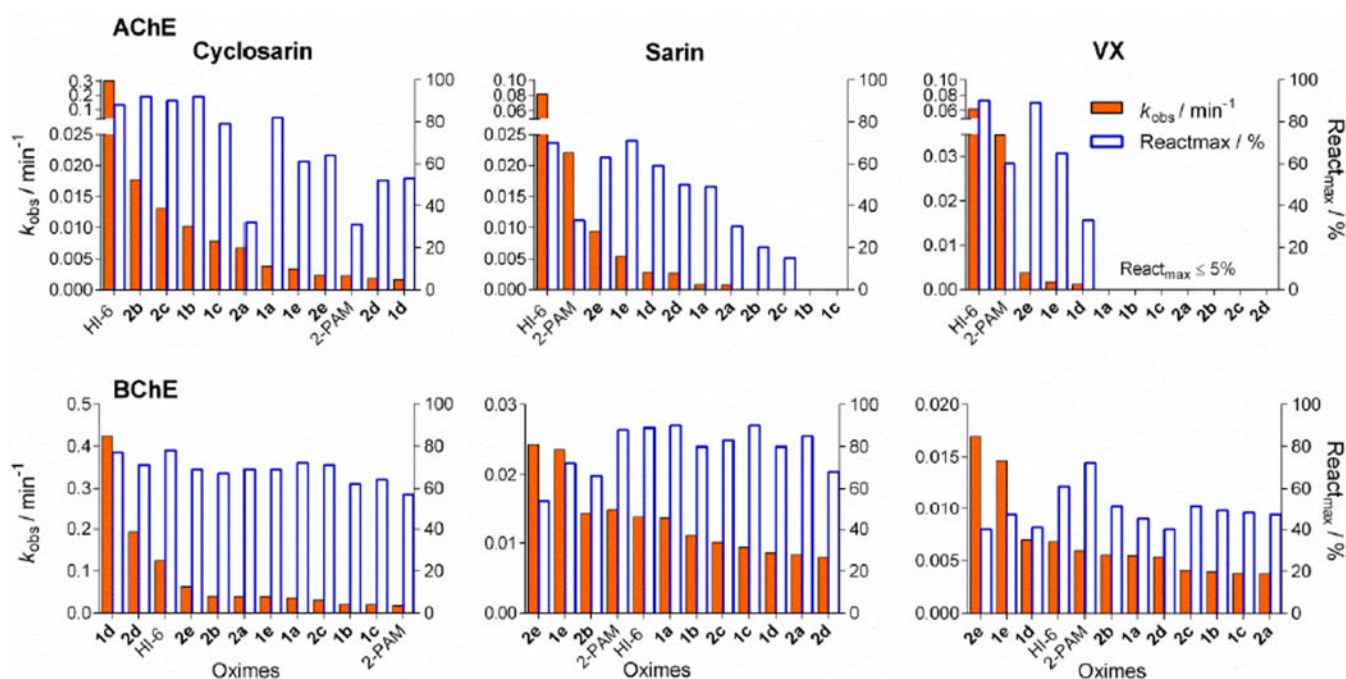


Figure 2.

Reactivation screening of cyclosarin-, sarin- and VX- inhibited human AChE and BChE with 10µM (AChE) or 100 µM (BChE) phenyltetrahydroisoquinoline aldoximes. The observed reactivation rates (k_{obs}) and maximal percentage of reactivation ($\text{React}_{\text{max}}$) within 24 h were evaluated and deviations were less than 10%. k_{obs} was not evaluated when $\text{React}_{\text{max}}$ was <30%. The results for tabun inhibition are not shown due to <5% reactivation within 24h.

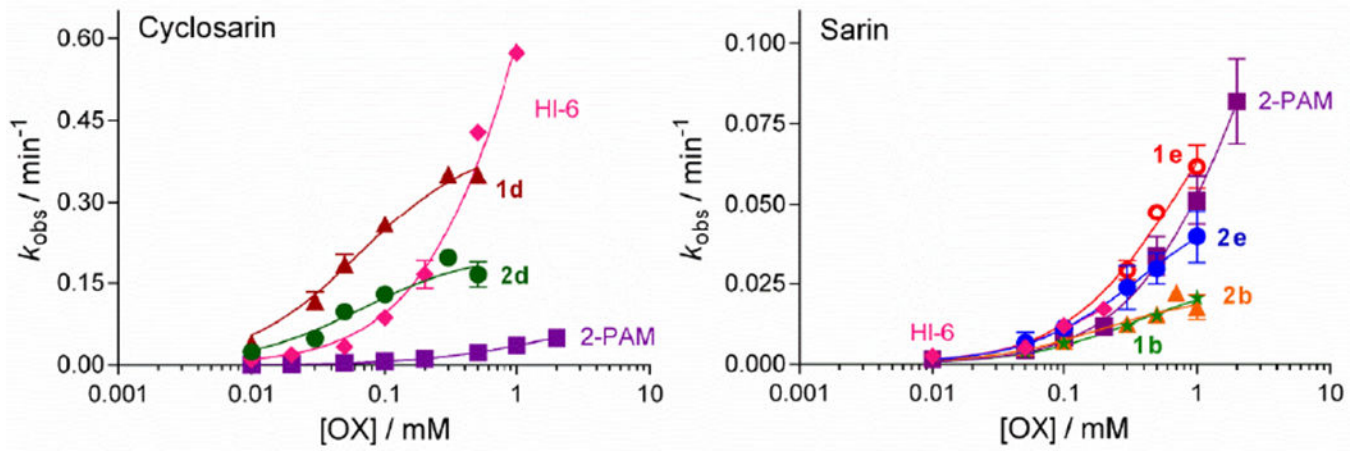


Figure 3.
Reactivation kinetics of cyclosarin- and sarin-inhibited BChE by the selected aldoximes.

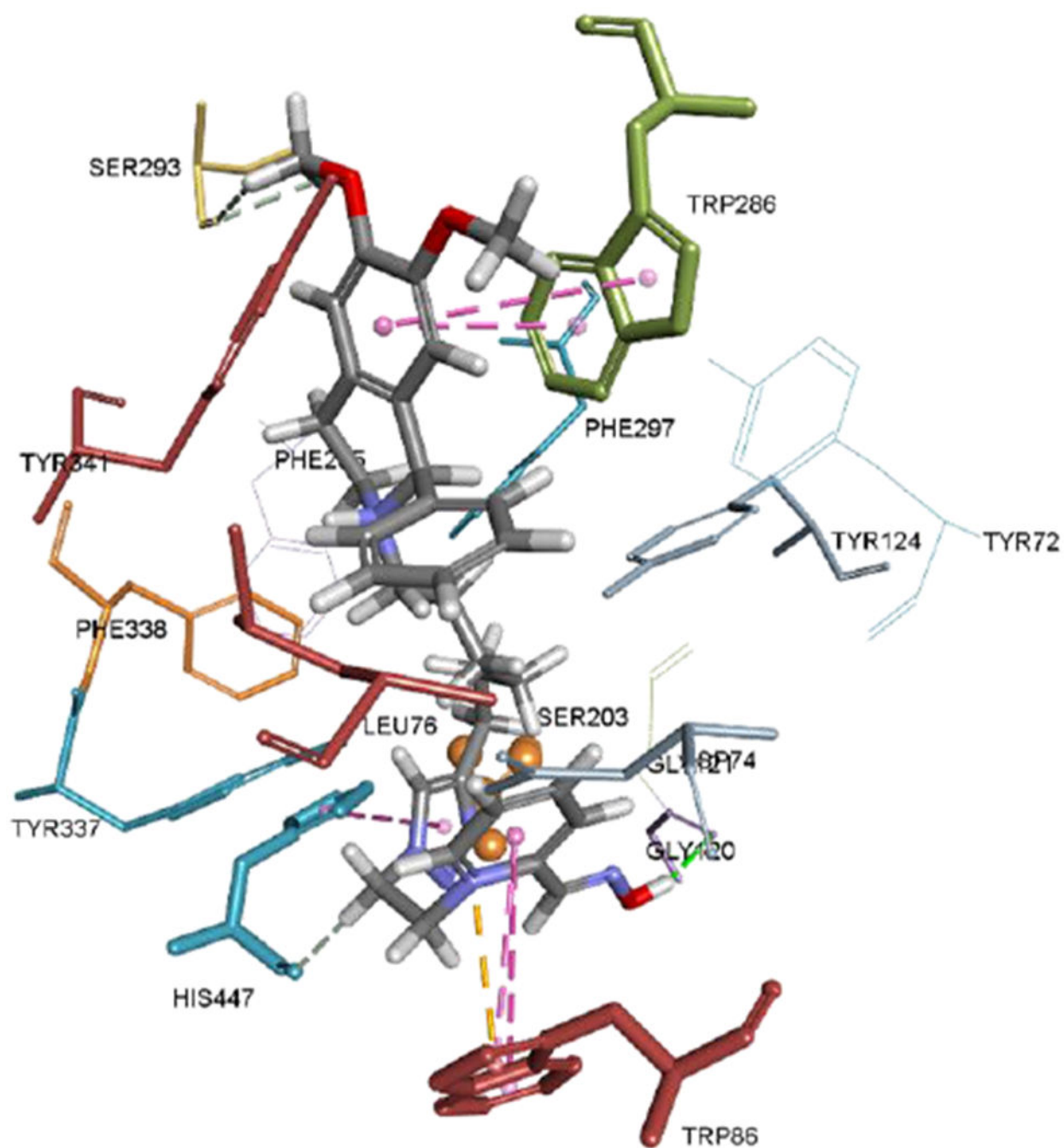


Figure 4. Positioning of the aldoxime **2a** with a protonated nitrogen of phenyltetrahydroisoquinoline ring in the mouse AChE active site (PDB code: 5EHN; Bourne et al, 2016). The catalytic Ser203 is presented as an orange stick and ball. Hydrogen bonds are shown in green, electrostatic interactions in orange and π - π interactions in magenta.

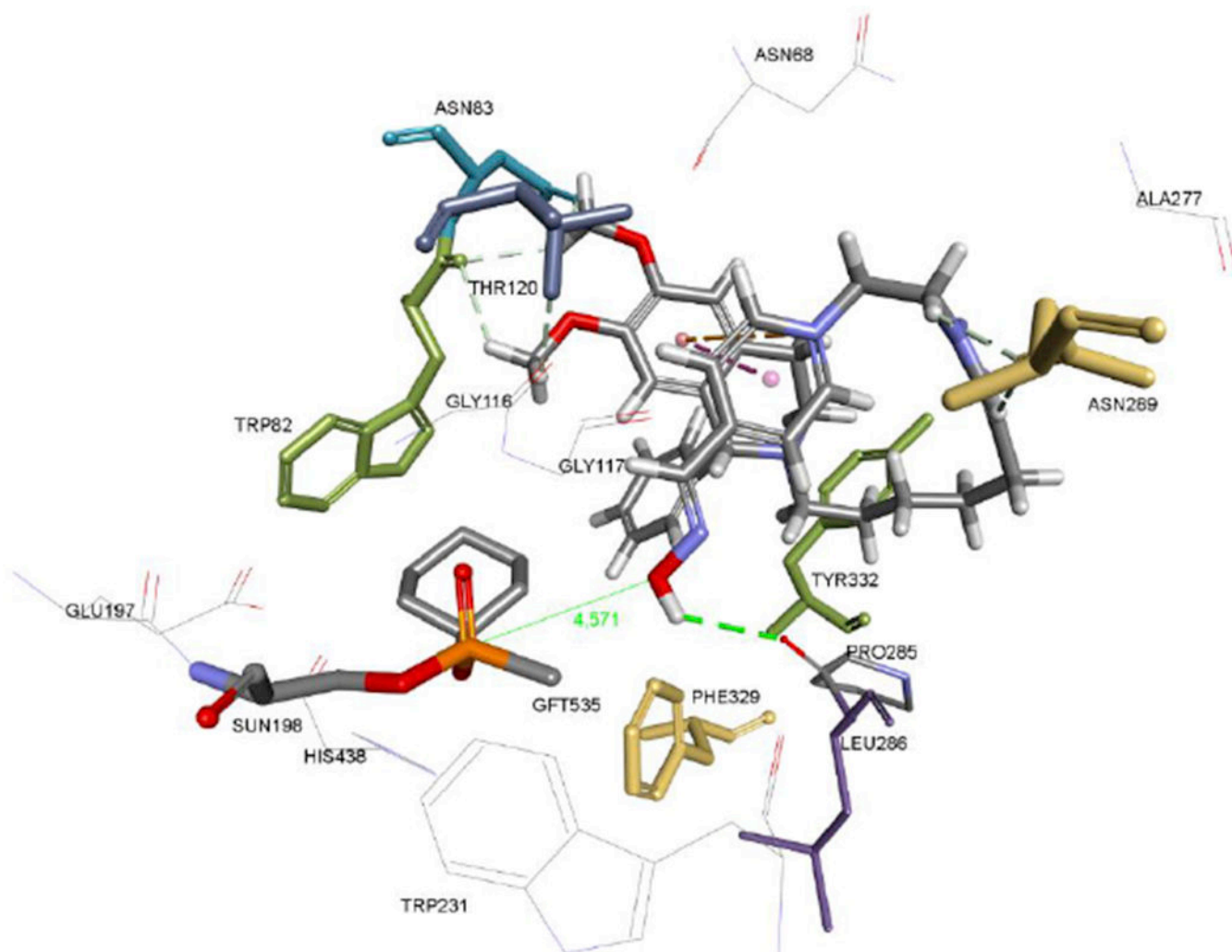
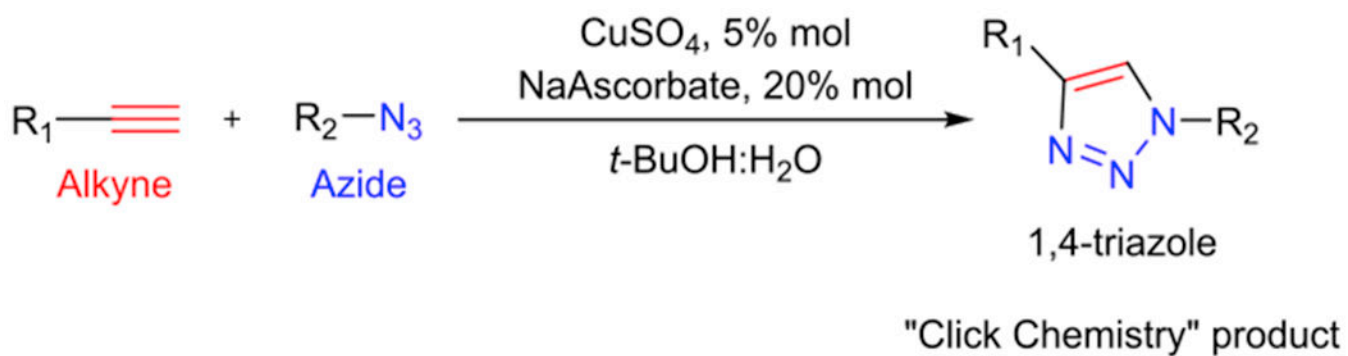


Figure 5. Minimized structure of aldoxime **1d** in cyclosarin-phosphonylated BChE modelled from human BChE inhibited by tabun (PDB code: 3DJY; Carletti et al, 2008). Residues Leu286, Phe329 and Tyr332 were set to be flexible during docking. Hydrogen bonds are shown in green, π - π and alkyl- π interactions in orange.

**Scheme 1.**

The copper(I)-catalyzed azide-alkyne cycloaddition employed to synthesize 1,4-triazole oximes.

Table 1.

Dissociation inhibition constants ($K_i \pm$ S.E.) of the phenyltetrahydroisoquinoline aldoximes for AChE and BChE determined at 25 °C.

Oxime	K_i (nM)		BChE/AChE
	AChE	BChE	
1a	15 ± 1	78 ± 7	5.2
1b	8.9 ± 0.5	8500 ± 1300	955
1c	67 ± 4	8600 ± 1700	128
1d	130 ± 8	4900 ± 500	38
1e	41 ± 4	10900 ± 1700	266
2a	6.0 ± 0.7	71 ± 4	12
2b	28 ± 3	5900 ± 500	211
2c	32 ± 3	4800 ± 700	150
2d	23 ± 2	147 ± 14	6.4
2e	100 ± 9	7900 ± 1200	79
HI-6 ^a	25000 ± 1400	215000 ± 11000	8.6
2-PAM ^a	180000 ± 10000	390000 ± 48000	2.2

Table 2.

Reactivation parameters determined for selected aldoximes and human BChE inhibited by cyclosarin and sarin. The results are expressed as mean \pm S.E. calculated from at least three experiments. Maximal percentage of reactivation ($\text{React}_{\text{max}}$) and the time (t) in which it was achieved were also determined.

Inhibitor	Oxime	k_2 / min^{-1}	$K_{\text{ox}} / \mu\text{M}$	$k_r / \text{M}^{-1} \text{min}^{-1}$	$\text{React}_{\text{max}} / \%$	t / min
Cyclosarin	1d	0.40 ± 0.02	66 ± 8.0	6250 ± 55	80	10
	2d	0.21 ± 0.02	67 ± 16	3050 ± 520	70	30
	HI-6	1.4 ± 0.27	1410 ± 390	1010 ± 95.0	90	30
	2-PAM	0.08 ± 0.01	1200 ± 240.0	65 ± 10	80	60
Sarin	1e	0.11 ± 0.013	800 ± 170	140 ± 14	80	60
	2e	0.058 ± 0.014	410 ± 250	140 ± 50	60	60
	2b	0.022 ± 0.0033	195 ± 89.0	110 ± 35	70	120
	1b	0.030 ± 0.003	365 ± 84.0	70 ± 10	90	180
	HI-6	0.041 ± 0.019	270 ± 200	150 ± 40	100	300
	2-PAM	0.18 ± 0.05	2400 ± 995.0	75 ± 10	100	120