

Combining Independent Drug Classes into Superior, Synergistically Acting Hybrid Molecules**

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Increasing the potency of synthesized drugs has been a stepwise process accomplished by progressively modifying the chemical scaffold of a single parent lead compound. To date, there has been no basis for thinking that the combination of pharmacological effects of independently acting drugs could be achieved beyond mere simultaneous administration. We reasoned that if the target molecule of two independent classes of drugs was the same, chemical synthesis of a hybrid compound where these drugs presented moieties within one molecule might yield synergistic effects; that is, a new quality might emerge that would be more than the sum of the single-

moiety compounds. Such multifunctional hybrid compounds that assign different functions to its different moieties to achieve a synergistic pharmacodynamic effect have successful predecessors in nature: for example, bleomycin is a natural compound with three different moieties acting in concert to cleave DNA.^[1]

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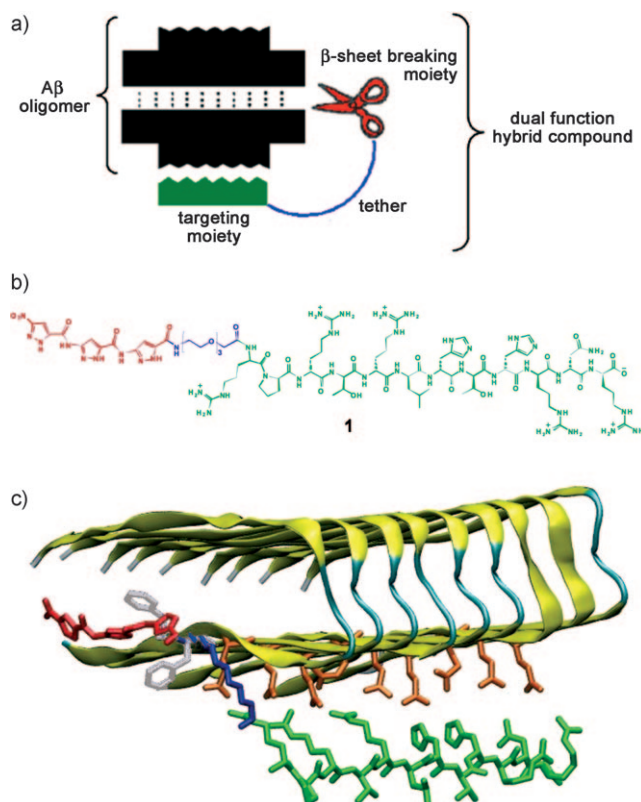


Figure 1. Design, function, and modeling of hybrid compounds.

a) Hybrid compound, composed of a recognition moiety docking on the target (here: A β oligomers) linked by a flexible tether to the functional moiety, a β -sheet breaker. b) Chemical structure of lead hybrid compound Trimer-TEG-D3 (**1**). c) Model of the A β /1 complex. The side chains of the glutamates (orange) in adjacent A β monomers (yellow) are located in close spatial proximity and are contacted by the basic residues of the D3 peptide (green). The aminopyrazole trimer (red) is bound to the A β backbone and interacts with the two phenylalanines (F19, F20; gray). The color coding of the individual parts of **1** is according to (a) and (b).

Herein we describe the design and chemical synthesis of hybrid compounds that target the disassembly of neurotoxic A β oligomers. Cross- β -sheeted A β oligomers or amyloids are a hallmark of Alzheimer's disease (AD). Misprocessing of an amyloidogenic fragment A β derived from proteolytic processing of amyloid precursor protein is critical in a cascade of events that starts with the oligomerization of A β and, ultimately, ends in neuronal death in the central nervous system.^[2] To date, rationally designed small-molecule β -sheet breakers, although highly efficient in preventing or disassembling β -sheet structures in cell-free *in vitro* systems, have failed to show convincing effects *in vivo* owing to their highly unspecific binding.

We reasoned that lack of *in vivo* efficiency of small β -sheet breakers might be overcome by the addition of a molecular recognition unit that would direct the β -sheet breaking moiety to its target molecule. Molecular recognition is a key property of polypeptides that can be identified in evolutionary algorithms, for example in phage display systems comprising iterative cycles of panning and selection of peptides. We identified D3, a D-enantiomeric dodecapeptide, as a potent A β oligomer binder in a mirror-image phage display,^[3,4] that is able to modulate A β aggregation, plaque load, and neuroinflammation processes in the brains of transgenic (APPswe/PS1 Δ E9) mice.^[3,5]

In our hybrid compound (Figure 1), we would thus combine two entirely different drug development strategies:

evolutionary selection from a peptide library together with rationally designed small molecules. As rationally designed β -sheet breakers, we chose aminopyrazoles (APs) that possess a specific donor–acceptor–donor (DAD) sequence of hydrogen bond donors and acceptors, that are perfectly complementary to that of a β sheet,^[6] binding selectively to the backbone of misfolded peptides residing in the β -sheet conformation,^[7] and able to disassemble preformed A β fibrils *in vitro*.^[8]

Comparative modeling and molecular dynamics simulations were used to investigate the interaction of A β ^[9] of AP linked to D3 with different linkers and to a pentalysine peptide (KKKKKG) as charge control for D3. The AP moiety of **1** was modeled to interact with the diaromatic motif F19/F20, and the D3 part was placed in spatial vicinity of a negatively charged surface patch of the A β protofibril to maximize electrostatic complementarity (see the Supporting Information, Figure S1). These simulations revealed that after 10 ns, the multiple interactions formed between D3 peptide and the E22 residues remained stable (Figure 1c; Supporting Information, Figure S2) whilst only one salt bridge was preserved between the pentalysine moiety and the A β fibril (Supporting Information, Figure S2D), thus providing an explanation for the enhanced activity of D3 compared to other substances (Figure 2a; Supporting Information, Figure S4B).

When added to cells that continuously secrete A β oligomer (7PA2 cells^[10]), the hybrid compound with the triethylen-

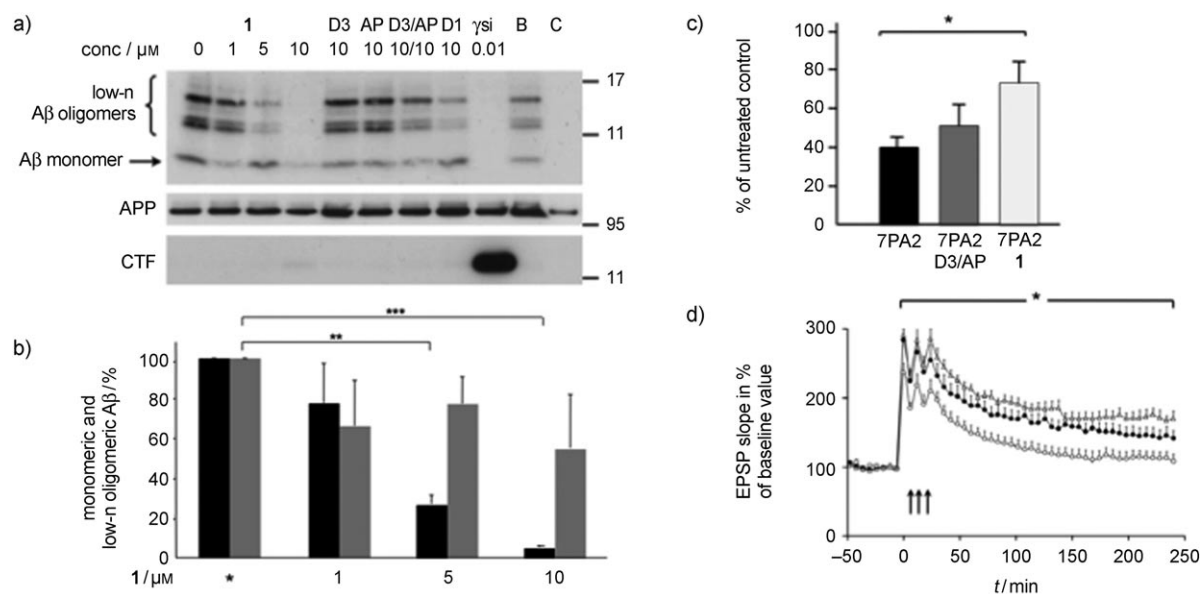


Figure 2. Biochemical and biological effects of hybrid compounds. a) Western blot of immunoprecipitated A β oligomers and monomers from 7PA2 cells that were treated with different substances. Addition of **1** led to a dose-dependent reduction of A β oligomers. Treatment with the D-peptide D3 or the aminopyrazole (AP) moiety alone, or as simultaneously applied substances (D3/AP) at 10 μ M, did not show any A β oligomer reducing effects in this assay. Further control experiments included a hybrid compound consisting of another D-peptide (10 μ M; D1), as well as γ -secretase inhibitor LY411575 (10 nM), 7PA2 cells treated with AP-dissolving substance DMSO at the same concentrations (B), and untransfected CHO cells not secreting A β oligomers (C). Expression of APP and APP-C-terminal fragments (CTF) as a control is shown in the lower panels, as indicated. b) Comparison of 5 independent experiments of 7PA2 cells treated with **1** at 1, 5, and 10 μ M, showing clear effects on oligomeric A β (*** p = 0.0007, ** p = 0.0082), whereas monomeric A β was not significantly affected (oligomeric A β black bars, monomeric A β gray bars). c) 10 μ M **1**, but not a composition of both single compounds at 10 μ M each, reversed the effect of 7PA2 supernatant on mEPSC frequency in cultured cortical mouse neurons. Significant reversal of mEPSC suppression (* p < 0.025 by Student's test after Bonferroni correction). Bars represent mean \pm SEM (see Supporting Information for details). d) Compound **1** prevented the A β -mediated decrease of LTP in acutely isolated hippocampal slices. Compared to control LTP (●), oligomeric A β (1–42) significantly reduced LTP (○), which was prevented by co-application of 10 μ M **1** (△) * p < 0.05 analysis of variance with repeated measures; significance between control and A β and between A β and **1**/A β .

glycol (TEG) linker, termed Trimer-TEG-D3 (**1**), inhibited A β oligomerization in a dose-dependent manner (Figure 2a,b). Remarkably, single compounds or the combination of both single-compound moieties administered at same concentrations did not produce any effect in this assay, thus demonstrating the dramatic synergistic action of both moieties within the composite compound that in effect leads to a new function (Figure 2a). Upon administration of **1**, only the decrease of the A β oligomer fraction was significant compared to the A β monomer fraction, and no increase of C-terminal APP fragments was observed, suggesting that **1** neither decreased A β monomer generation nor inhibited γ secretase (Figure 2a). Even very high concentrations of each single compound or their combination did not lead to a reduction in A β oligomers (Supporting Information, Figure S3), and **1** revealed cytotoxicity only at concentrations of more than 100 μ M (Supporting Information, Figure S4).

To demonstrate the specificity of the D3 moiety in targeting A β oligomers, a hybrid with a D-peptide that recognizes a different epitope of A β , termed D1, was also tested.^[4] With this compound, only an insignificantly weak inhibition of A β oligomerization was observed (Figure 2a). Moreover, as predicted from molecular modeling, the AP-pentylsine hybrid had no effect on A β oligomer assembly (Supporting Information, Figure S5B).

Our concept of synergistic hybrid compounds predicts a critical importance of linker length between moieties. Structural and energetic considerations suggest that a shorter linker should favor a tight interaction owing to the smaller entropic loss upon binding, whereas longer linkers should be unfavorable. When we synthesized hybrids with no spacer, a shorter γ -aminobutyric acid (GABA) spacer, or a longer (TEG)₂ spacer (Supporting Information, Figure S5A), we verified this relation in that only no spacer or a five-atom linker (GABA) showed markedly increased A β oligomer inhibition (Supporting Information, Figure S5B,C).

Synaptic pathology is a key biological effect of A β oligomers, which should be reversed by **1**. Compound **1** blocked the A β -induced decrease in mEPSC frequency (mEPSC = miniature excitatory post-synaptic current) that is mediated by AMPA receptors^[11] in cultured cortical neurons (Figure 2c), but it did not affect the mean mEPSC amplitude. The effect was achieved with 10 μ M **1** but not with the single compounds or a combination of both single compounds (Figure 2c). Similarly, A β -oligomer-induced impairment of long-term potentiation (LTP),^[12] a form of synaptic plasticity in acute hippocampal slices, could be reversed by coapplication of 10 μ M **1** (Figure 2d). These results demonstrated that **1** prevented A β -induced synaptotoxicity in two independent assays.

Multimer growth to amyloid fibrils should be inhibited by increased binding of the two moieties of **1** to A β oligomers. Analytical ultracentrifugation of synthetic A β in vitro^[13] showed an equal distribution across all multimeric fractions (Figure 3a, upper panel). When preincubated with **1**, A β partitioned to high-molecular-weight fractions at the expense of oligomeric fractions (Figure 3a, middle panel), as evidence that direct binding of **1** to A β oligomers influenced A β oligomer assembly. As A β thioflavin T (ThT) fluorescence

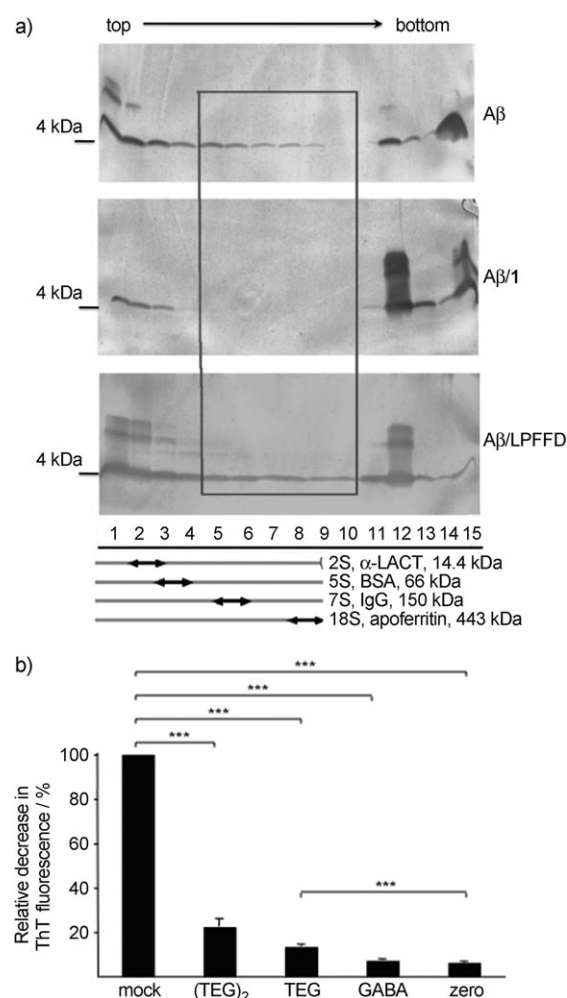


Figure 3. Biophysical characterization of the effect of hybrid compounds on A β assembly. a) Analysis of the effect of **1** on A β aggregation by analytical density gradient ultracentrifugation analyzed by SDS-PAGE and silver staining. Continuous fractions from light (left) to heavy (right) were loaded: monomeric A β in the leftmost fractions (lane 1), oligomeric A β in the fractions in the red box (lanes 5–10), and multimeric aggregated A β in fractions 12 and higher. Compared to untreated A β (125 μ M; top panel), or control-treated A β (bottom panel), compound **1** (62.5 μ M; middle panel) led to a shift of oligomeric fractions to high-molecular-weight, non-amyloid complexes. Calibration of fractions for sedimentation coefficients 5 (black double arrows) with α -lactalbumin (α -LACT), bovine serum albumin (BSA), immunoglobulin G (IgG), and apoferritin. b) Inhibition of ThT-positive A β fibrillogenesis by **1** (TEG) and hybrid compounds with shorter (GABA) or longer ((TEG)₂) linker length or without linker (zero). (mean \pm standard deviations of results, four separate runs, three replicates per run; *** $p \leq 0.001$ by Student's *t*-test). Clear anti-amyloid effects of all hybrid compounds are observed, with the zero linker being strongest.

(an indicator of amyloid fibril content) decreased upon application of **1** (Figure 3b), these higher fractions could not represent A β amyloid fibrils but rather irregularly structured high-molecular-weight complexes. Accordingly, transmission electron microscopy revealed the total lack of any remaining fibrillar structures in A β /1 mixtures (Supporting Information, Figure S6), thus demonstrating that **1** inhibited A β oligomer assembly by driving A β aggregation

to nontoxic, non-amyloid complexes. As for native A β oligomer assembly inhibition in 7A2 cells (Supporting Information, Figure S5C), we also observed a structure–activity relationship of hybrid compounds in the cell-free ThT assay (Figure 3b), confirming that the zero-linker compound has strongest activity. Our biophysical analysis by three independent methods (density gradient centrifugation, ThT assay, and transmission electron microscopy), as well as our findings that monomeric A β is not elevated when A β oligomers are decreased (Figure 2b, 3a), lead us to conclude that rather than inhibiting A β oligomer assembly, hybrid compounds lead to incorrect, bioinactive, and non-amyloid A β misassembly, which could eventually be degraded more easily.

In conclusion, we demonstrated that chemical synthesis of two entirely different substance classes acting on the same target can be covalently linked to yield dramatic synergistic effects and lead to novel properties. Taking A β oligomers as an example, we also showed that two entirely different principles of drug development can be combined: polypeptides developed through mirror image phage display, which takes advantage of evolutionary algorithms to select for molecular recognition, and the rationally designed small-molecule β -sheet breakers. Our investigations may spark similar efforts with different target molecules and thereby greatly accelerate drug development.

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- [1] B. Meunier, *Acc. Chem. Res.* **2008**, *41*, 69–77.
- [2] C. Haass, D. J. Selkoe, *Nat. Rev. Mol. Cell Biol.* **2007**, *8*, 101–112.
- [3] T. van Groen, K. Wiesehan, S. A. Funke, I. Kadish, L. Nagel-Steger, D. Willbold, *ChemMedChem* **2008**, *3*, 1848–1852.
- [4] K. Wiesehan, K. Buder, R. P. Linke, S. Patt, M. Stoldt, E. Unger, B. Schmitt, E. Bucci, D. Willbold, *ChemBioChem* **2003**, *4*, 748–753.
- [5] S. A. Funke, T. van Groen, I. Kadish, D. Bartnik, L. Nagel-Steger, O. Brener, T. Sehl, R. Batra-Safferling, C. Moriscot, G. Schoehn, A. H. C. Horn, A. Müller-Schiffmann, C. Korth, H. Sticht, D. Willbold, *ACS Chem. Neurosci.* **2010**, *1*, 639–648.
- [6] T. Schrader, C. Kirsten, *J. Am. Chem. Soc.* **1997**, *119*, 12061–12068.
- [7] P. Rzepecki, T. Schrader, *J. Am. Chem. Soc.* **2005**, *127*, 3016–3025.
- [8] P. Rzepecki, L. Nagel-Steger, S. Feuerstein, U. Linne, O. Molt, R. Zadnarm, K. Aschermann, M. Wehner, T. Schrader, D. Riesner, *J. Biol. Chem.* **2004**, *279*, 47497–47505.
- [9] T. Lührs, C. Ritter, M. Adrian, D. Riek-Loher, B. Bohrmann, H. Dobeli, D. Schubert, R. Riek, *Proc. Natl. Acad. Sci. USA* **2005**, *102*, 17342–17347.
- [10] M. B. Podlisny, B. L. Ostaszewski, S. L. Squazzo, E. H. Koo, R. E. Rydell, D. B. Teplow, D. J. Selkoe, *J. Biol. Chem.* **1995**, *270*, 9564–9570.
- [11] a) G. M. Shankar, B. L. Bloodgood, M. Townsend, D. M. Walsh, D. J. Selkoe, B. L. Sabatini, *J. Neurosci.* **2007**, *27*, 2866–2875; b) G. M. Shankar, S. Li, T. H. Mehta, A. Garcia-Munoz, N. E. et al., *Nat. Med.* **2008**, *14*, 837–842.
- [12] M. J. Rowan, I. Klyubin, Q. Wang, N. W. Hu, R. Anwyl, *Biochem. Soc. Trans.* **2007**, *35*, 1219–1223.
- [13] R. V. Ward, K. H. Jennings, R. Jepras, W. Neville, D. E. Owen, J. Hawkins, G. Christie, J. B. Davis, A. George, E. H. Karran, D. R. Howlett, *Biochem. J.* **2000**, *348*, 137–144.