

Review

Comprehensive Survey of Combinatorial Library Synthesis: 2001

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Reviews

Comprehensive Survey of Combinatorial Library Synthesis: 2001

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This is the fifth review in an annual comprehensive survey series on combinatorial chemistry.¹ There were a total of 305 small-molecule libraries published in 2001 as delineated in Tables 1–10. The format remains the same as in previous reviews with libraries divided into two broad categories: libraries with disclosed biological activity (78 entries; Tables 1–5) and those without associated biological screening data (227 entries; Tables 6–10). Two new tables were added this year, recording solid-phase reagents and scavengers (Table 11) and solid-phase linkers (Table 12) reported in 2001.

The percentage of academic research groups (60%) publishing solid- and solution-phase research continued to outpace industrial groups by a moderate margin. Three years ago, the opposite was observed. Academic groups are the principal drivers for new solid-phase methodology and the synthesis of libraries based on natural-product templates.

Although the number of published libraries synthesized on solid-phase continues to dominate, solution-phase library synthesis, which includes the use of resin-bound reagents and scavengers, is clearly on the rise. Some 60% of the libraries found in Tables 1–5 were prepared using solutionphase chemistry, up over 150% from the previous year. Successful optimization programs in 2001 that relied on solution-phase chemistries include cathepsin K inhibitors,²²⁸ raf kinase inhibitors,²⁷⁴ Cdk4 inhibitors,²⁷⁴ and CCR3 antagonists.²⁷⁴ In contrast to libraries published with screening results, only 22% of the methodology-type libraries (Tables 6–10) were solution-phase.

The average size of the libraries in Tables 1-5, i.e., the number of compounds per library, decreased in 2001 (50%

contain <100 members; 86% contain <1000 members). Several years ago, it was common to see the synthesis and screening of large libraries containing thousands of members. The current trend is to apply combinatorics in the iterative process of lead optimization by quickly creating small sets of libraries to address specific SAR and pharmacokinetic queries. Pharmaceutical companies appear to be outsourcing to an increasingly greater extent the synthesis of general discovery libraries. For example, Merck, Pfizer, Roche, and Pharmacia signed multivear collaboration agreements in 2001 with companies such as Discovery Partners International and Evotec OAI to synthesize small-molecule chemical libraries. Some major pharmaceutical firms have reportedly scaled back their dedicated combinatorial chemistry departments, redirecting resources to medicinal chemistry optimization. Academic institutions, on the other hand, are creating new departments, schools, and centers of excellence for streamlined synthesis.

Selected library highlights for the year include an inspiring account of the discovery and optimization of selective Cdk4 inhibitors via an integrated strategy of computational lead generation, homology modeling, structure-based design, library synthesis, and traditional analogue synthesis.^{109,110} The strength of combinatorial chemistry to simultaneously conduct "multiple-point modifications" was aptly demonstrated by Smith and co-workers²²⁷ in a "hit-to-lead" raf kinase inhibitor campaign. Combinatorial chemistry allowed the Smith group to break free of a SAR paradigm defined by "sequential fragment-based" optimization, ultimately leading to a clinical candidate. The Lead Discovery group at Organon

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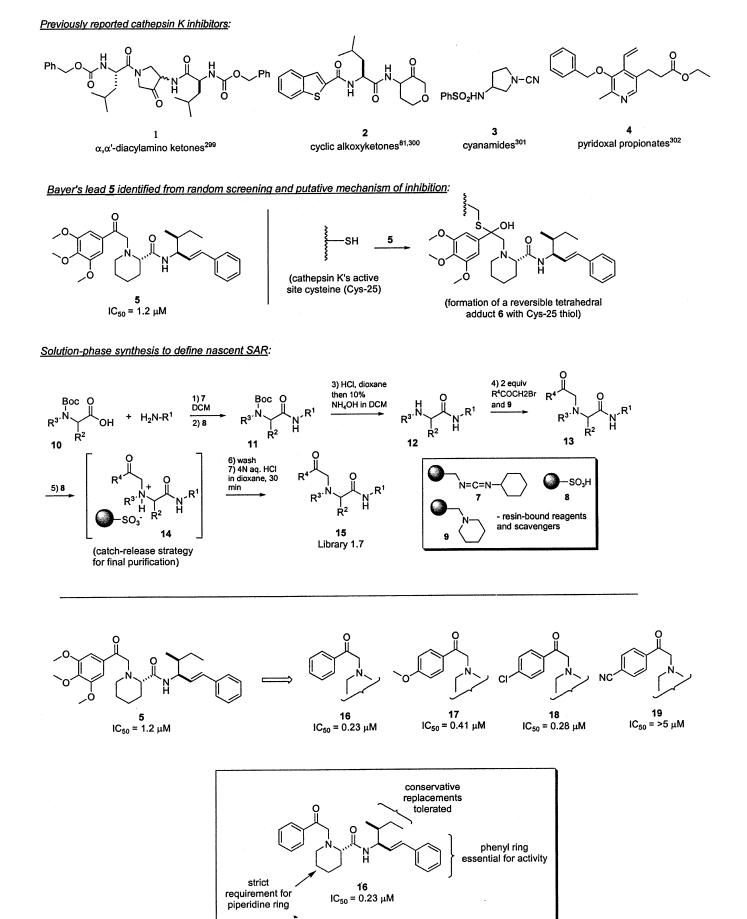


Figure 1. Aminomethyl ketones as cathepsin K inhibitors.²²⁸

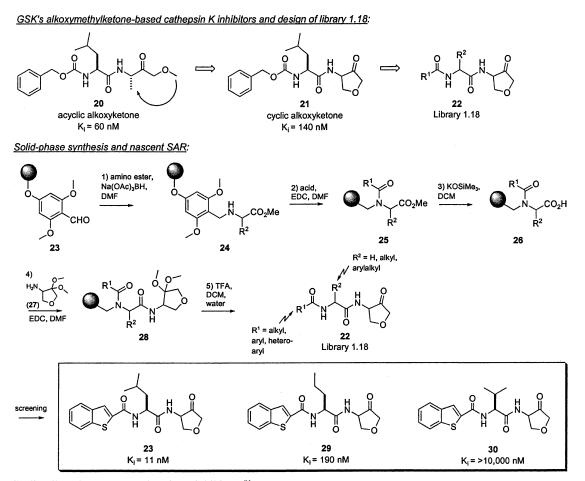


Figure 2. Cyclic alkoxyketones as cathepsin K inhibitors.⁸¹

provided a detailed design of a general screening library biased toward central nervous system (CNS) bioavailability.^{20,49} Finally, GlaxoSmithKline (GSK) described a thwarted account of optimizing a structurally complex oxytocin antagonist that was obtained as a screening hit from a combinatorial library.^{80,274} Their work highlights the difficulty in simplifying complex, high molecular weight leads to optimize potency, selectivity, and pharmacokinetic parameters. The GSK report supports the contentions of Teague,³¹⁵ Hann,³¹⁶ and Oprea³¹⁷ who argue for library designs that yield simpler, low molecular weight structures as starting points for optimization. Detailed descriptions of these and other highlights are given below.

1. Biologically Active Libraries (Tables 1–5)

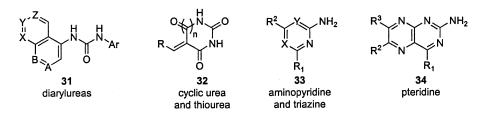
Cathepsin K Inhibitors.^{81,228} Cathepsin K is a cysteine protease selectively expressed in osteoclasts and has been implicated in bone resorption. Inhibitors of the enzyme have been shown to reduce bone loss in vivo, supporting the hypothesis that such agents may be clinically useful therapeutics for the treatment of diseases characterized by excessive bone loss, e.g., osteoporosis. Several structurally distinct classes of compounds have been identified as cathepsin K inhibitors including α, α' -diacylamino ketones 1, acyclic³⁰⁰ and cyclic alkoxyketones 2,⁸¹ cyanamides 3,³⁰¹ and pyridoxal propionate derivatives 4³⁰² (Figure 1). During a high-throughput screening campaign, Smith and co-workers

at Bayer Research discovered α -amino aryl ketones as a new class of cathepsin K inhibitors.²²⁸ Screening hit 5, $IC_{50} =$ 1.2 μ M, was thought to be a reversible inhibitor by virtue of its potential to form covalent tetrahedral adduct 6 with the active-site Cys-25. To explore the SAR of the aryl ketone fragment in 5, an expedient solution-phase synthesis of library 1.7 (15) was developed, facilitated by polymersupported reagents and scavengers. The synthesis began with coupling N-Boc-protected amino acids 10 to both custom and commercially available amines using resin-bound carbodiimide 7 as the activating reagent. Excess amines were scavenged with resin-bound sulfonic acid 8. After removal of the N-Boc protecting group with HCl/dioxane, the resulting amine HCl salts were free-based with 10% NH₄OH/ DCM to furnish amines 12. Alkylation of 12 was carried out by treatment with 2 equiv of an α -halomethyl ketone in the presence of resin-bound piperidine 9. Final purification of library members was achieved by a "catch-and-release" strategy in which products were first captured as their resin-bound sulfonic acid salts 14, followed by thorough resin washing, and then liberated from resin upon exposure to 4 N aqueous HCl in dioxane. Evaluation of compound arrays against cathepsin K revealed a range of inhibitory activities in which less bulky but electron-rich or lipophilic aryl groups yielded more potent inhibitors, e.g., 16, $IC_{50} = 0.23 \ \mu M$. Additional analogues were subsequently prepared to explore other regions of 5 to define the nascent SAR.

Structure-based approach:

- Created homology model of Cdk4 from X-ray structure of Cdk2
- de novo design program LEGEND generated novel templates from homology model
- In-house SEEDS program then sifted through substructures of LEGEND outputs to identify
- commercially available or synthetic feasible derivatives

- Computational exercise led to four structural templates 31-34 possessing 15-500 μ M affinity for Cdk4



After selecting diraryl ureas (e.g., 35) as most tractable class, "informer" libraries 2.2 and 2.3 were synthesized:

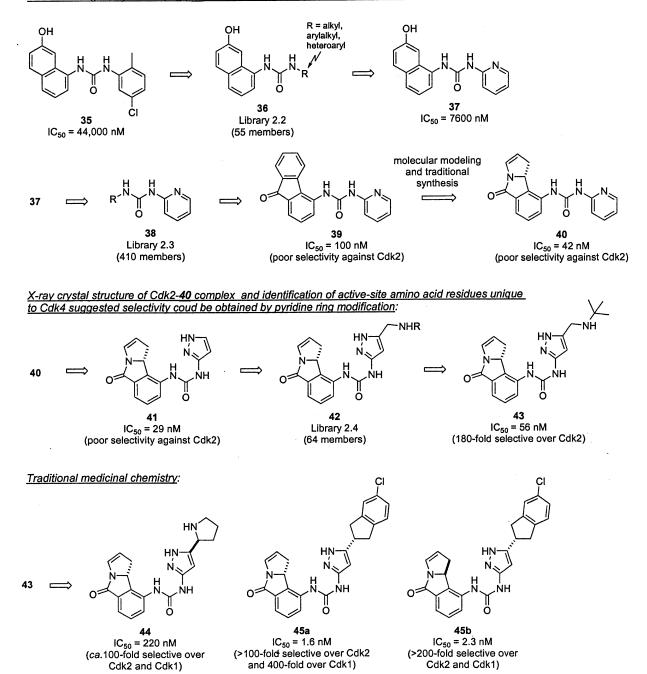


Figure 3. Development of selective Cdk4 kinase inhibitors.^{109,110}

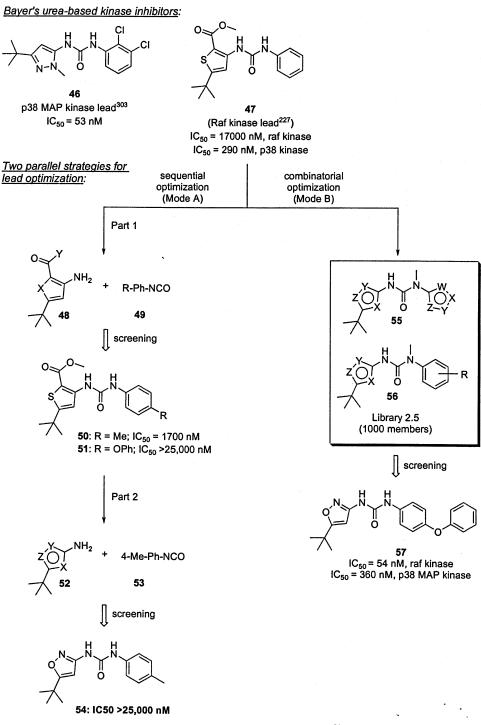
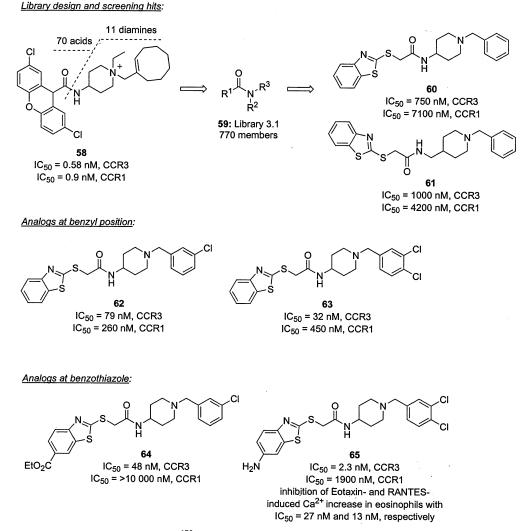
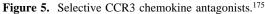


Figure 4. Library 2.5 yielding second-generation urea-based Raf kinase inhibitor lead.⁸¹

As part of an ongoing cathepsin K inhibitor program, researchers at GlaxoSmithKline described the solid-phase preparation of library 1.18 (**22**) based on cyclic alkoxymethyl ketone template **22** (Figure 2).⁸¹ This library was designed to explore the SAR of initial lead **21** ($K_i = 140$ nM, cathepsin K), an inhibitor that was some 2-fold less active than its acyclic counterpart **20**. The synthesis commenced with the reductive amination of amino esters (10 equiv of each amine and Na(OAc)₃BH, DMF) and aldehyde resin **23**. The resin-bound secondary amines **24** were then acylated with a variety of carboxylic acids (3 equiv) employing standard coupling conditions. Hydrolysis of the ester with aqueous base led to racemization at R² in **26**. This

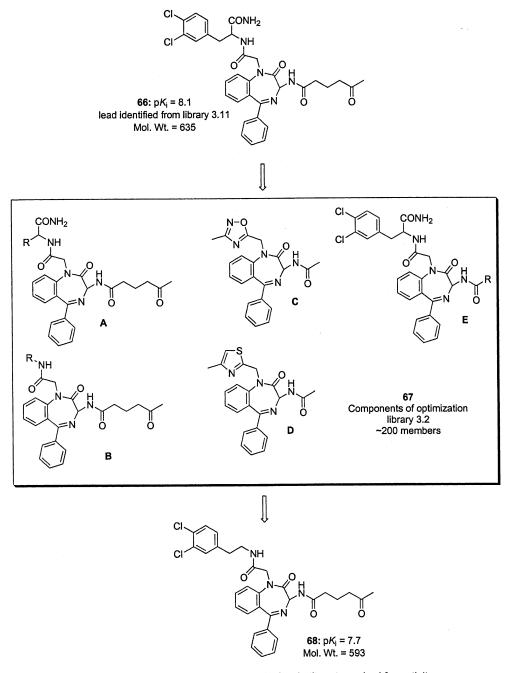
problem was circumvented by treatment with potassium trimethylsilanoate (10 equiv) in DCM to furnish chiral acid **26**, which in turn was coupled to the protected keto amine **27** yielding **28**. Library 1.18 was obtained upon acid treatment (TFA/DCM/H₂O (7:2:1)) of resin **28**, simultaneously liberating the salient ketone functionality. The synthesis was semiautomated via application of the ACT496 robotics or encoding with IRORI radiofrequency tags. The most potent compound identified from the library was cyclic ketone **23** ($K_i = 11$ nM), some 10-fold more potent than lead **21**. On the basis of the poor affinity observed for analogues **29** and **30**, the leucyl side chain in **23** appeared to be optimal.





Cyclin-Dependent Kinase 4 (Cdk4) Inhibitors.^{109,110} A new class of highly selective Cdk4 inhibitors was identified by Honma and co-workers using an integrated drug discovery strategy of structure-based design, library synthesis, and traditional analogue synthesis (Figure 3).^{109,110} The first step was the construction of a Cdk4 homology model derived from the crystal structure of Cdk2.109 This was accomplished using the de novo design program LEGEND, which sequentially generated some 1000 structures inside the deep, narrow cavity of Cdk4's ATP binding site. Typical of outputs from de novo design programs, many of the putative inhibitors produced by LEGEND were neither commercially available nor synthetically feasible. To overcome this issue, a second in-house computational program "SEEDS" was used to dissect the structures to afford pharmacophore queries. The queries in turn were used to survey compound databases. For instance, on searching the Available Chemicals Directory (ACD), some 4884 compounds (MW < 350) were culled, from which ca. 380 compounds were purchased. When screened in a cyclin-Cdk4 assay, 18 hits were obtained with IC_{50} values of 15-500 μ M. The hits were classified into four structural types 31–34. Urea 35 (IC₅₀ = 44 μ M), belonging to the class of diarylureas 31, was designated as the lead for SAR development. Two complementary "informer" libraries 2.2 (36) and 2.3 (38) based on 35 were sequentially synthesized using solution-phase methods (465 compounds total). Urea 39 (IC₅₀ = 100 nM) emerged, possessing a 440-fold improvement in binding affinity over lead 35. Molecular modeling and traditional synthesis led to urea 40 (IC₅₀ = 42 nM). An X-ray analysis of the Cdk2-40 complex was obtained, confirming the predicted binding mode in the ATP binding site.

Although urea 42 was reasonably potent against Cdk4, it lacked selectivity against Cdk2 and related kinases. Examination of the Cdk2-40 crystal structure and identification of active-site residues unique to Cdk4 suggested that modification of the pyridine ring in 40 might lead to more selective inhibitors.¹¹⁰ Exchange of the pyridine for a pyrazole ring afforded inhibitor **41** (IC₅₀ = 29 nM). On the basis of X-ray crystallographic studies and homology modeling, it was thought that introducing a C(5) substituent into the pyrazole heterocycle would lead to further enhancement in potency and selectivity. Hence, library 2.4 (42; 64 members) was prepared, yielding 43 as a potent inhibitor of Cdk4 with 180-fold selectivity over Cdk2. Further analogue synthesis furnished ureas 45a and 45b, $IC_{50} = 1.6$ and 2.3 nM against Cdk4, respectively, both possessing high selectivity across a panel of kinases. Cell-based assays demon-



- substituted benzyl and 3-(5-oxo-hexano-amide)-substituent required for activity

- 68 showed improved pharmacokinetics but less active than 66

- highlights difficulties experienced when optimizing complex structures

- underscores preferred library designs of less complex, lower MW structures

Figure 6. Attempted optimization of oxytocin antagonist 66.^{80,274}

strated that **45b** caused G_1 arrest in Rb(+) cancer cell line (Molt-4).

Raf Kinase Inhibitors.²²⁷ One clinically promising strategy for the treatment of cancer is the inhibition of raf kinase. This kinase is a downstream effector of ras, a signal transduction pathway involved in the regulation of cell differentiation and division. Activating mutations of ras are found in nearly one-third of all human cancers. Several years ago, Smith and co-workers at Bayer discovered heteroaryl ureas, e.g., **46**, as potent inhibitors of p38 MAP kinase (Figure 4).³⁰² This group found that this compound class also inhibits raf kinase, suggesting that heterocylic ureas may

represent a privileged kinase inhibitor motif.²²⁷ Lead **47** emerged from a high-throughput screening campaign. It has an IC₅₀ = 17 μ M against raf kinase, although it is a much more potent inhibitor of p38 MAP kinase, IC₅₀ = 0.29 μ M. Two parallel approaches were taken to improve the potency and selectivity of **47** for raf kinase. These included a "sequential" optimization strategy (mode A) in which the thienylamine was held constant while the aniline unit was varied (**48** + **49**) and vice versa (**52** + **53**) and a "combinatorial library" optimization strategy (mode B) in which broad combinations of heterocycles and anilines (library 2.5; **55** and **56**) were synthesized. Despite extensive analoguing

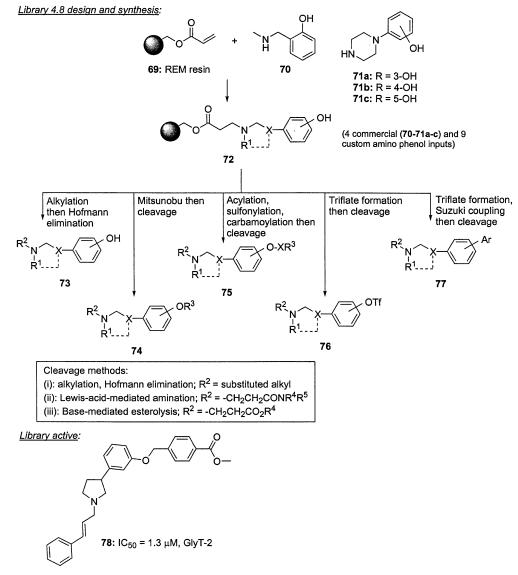


Figure 7. Library 4.8 synthesized using REM resin methodolgy.²⁰

Table 13. Molecular Property Anal	ysi	is
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molecular property	66% of CNS-active drugs (displayed property range)	library 4.8 (% members with property)	GlyT-2 inhibitor 78
molecular weight	125-425	$98\% \le 450$	427
CLogP	1.5-6.5	$80\% \le 6$	6.24
no. of H-bond donors	0-2	$99\% \le 1$	0
no. of H-bond acceptors	1-2	$40\% \le 2 (75\% \le 3)$	4
no. of rotatable bonds	0-7	$55\% \leq 7(80\% \leq 9)$	9
cLogBB ^{20b}	-0.74 to 0.77	$75\% \le 0.75 \ (98\% \le 1)$	0.5

via the sequential mode A, raf kinase potency could not be improved beyond the 1 μ M level. For example, introducing a *p*-methyl group in the phenyl ring of **47** gave urea **50** (IC₅₀ = 1.7 μ M; a 10-fold increase in binding), while many other analogues including **51** and **54** were inactive (>25 μ M). In contrast, compound **57**, with a raf kinase IC₅₀ = 54 nM and showing 7-fold selectivity versus p38 MAP kinase, was identified from the combinatorial library mode B. Urea **57** clearly lies outside the SAR established by sequential optimization. On the basis of the inhibitory activities of **51** and **54**, compound **57** would have been dismissed as an "inactive analogue" and therefore would not have been prepared in mode A optimization. This exercise demonstrates the power of combinatorial chemistry to conduct simultaneous "multiple-point modifications" to identify novel biologically active agents. Further studies on this class of compounds led to the identification of a clinical candidate.²²⁷

Chemokine Receptor CCR3 Antagonists.¹⁷⁵ Activation of the CCR3 receptor is an important determinant in the recruitment and accumulation of eosinophils to inflammatory sites characteristic of allergic diseases such as asthma. Thus, antagonists of CCR3, expected to suppress eosinophil infiltration, may have clinical potential to ameliorate allergic diseases. Banyu Tsukuba Research Institute reported amide **58** as a subnanomolar dual antagonist of CCR1 and CCR3

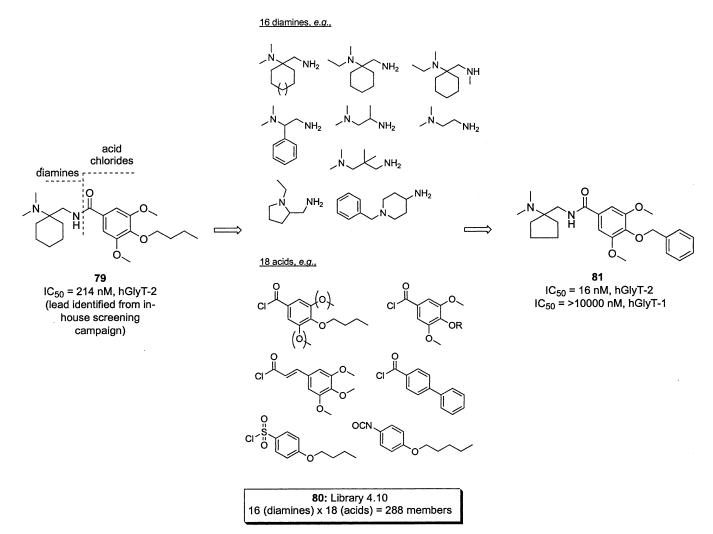


Figure 8. Glycine transporter type-2 inhibitors.⁴⁹

(Figure 5). Preliminary SAR studies of 58 demonstrated the importance of the carbonyl group as a hydrogen bond acceptor, a basic amine for electrostatic interaction to CCR3, and hydrophobic groups flanking the amide carbonyl and basic amine. It was also recognized that the quaternary ammonium group in 58 was a barrier to oral absorption and hence must be removed. Using this information, amide library 3.1 (59) was designed and synthesized using 70 carboxylic acid and 11 amine inputs, generating 770 total compounds. Inputs were culled from commercial and inhouse collections after clustering based on the binary Tanimoto coefficient and calculated by the MDL keys as a structural descriptor. Screening of the library furnished only two active compounds 60 and 61 possessing micromolar affinity for CCR3 and 4- to 10-fold selectivity over CCR1. Although the 1000-fold loss in receptor potency relative to 58 was disappointing, 60 and 61 represented secondgeneration leads that no longer contained the quaternary nitrogen. Amide 61 proved to be most amenable to SAR development. Sequential optimization, first at the benzyl moiety and then at the benzothiazole ring, gave rise to substantial improvements in affinity and potency. In particular, introducing chlorine atoms onto the benzyl ring increased CCR3 receptor affinity 50- to 100-fold, with IC₅₀ values of 79 and 32 nM for 62 and 63, respectively.

Incorporation of an amino group in the benzothiazole ring furnished **65**, a highly potent and selective antagonist: $IC_{50} = 2.3$ nM versus CCR3; $IC_{50} = 1900$ nM versus CCR1. Amide **65** inhibited Eotaxin- and RANTES-induced Ca²⁺ increase in eosinophils. It is interesting to note that if the benzothiazolylacetic acid had not been one of the 70 carboxylic acid library inputs, neither amides **60** and **61** would have been discovered nor subsequent SAR studies carried out leading to **65**.

Oxytocin Antagonists.^{80,274} The potent oxytocin antagonist 66 was identified at GlaxoSmithKline directly from a fully encoded differential release library (Figure 6).80 Compound **66** ($pK_i = 8.1$) displayed a poor pharmacokinetic profile, and an optimization program was undertaken to enhance both oxytocin antagonism and pharmacokinetic parameters. Given the high molecular weight and number of sites for metabolism, library design and SAR investigations centered on reducing the molecular weight and structural complexity of 66. Compound arrays of more than 200 new analogues were prepared using a combination of solid- and solution-phase synthesis (library 3.2). The first library members were modifications to the 3,4-dichlorobenzyl substituent (67A). Replacement of the phenyl ring with alkyl or cycloalkyl groups and removal of the chlorine atoms or their replacement with fluorine all resulted in loss of binding affinity.

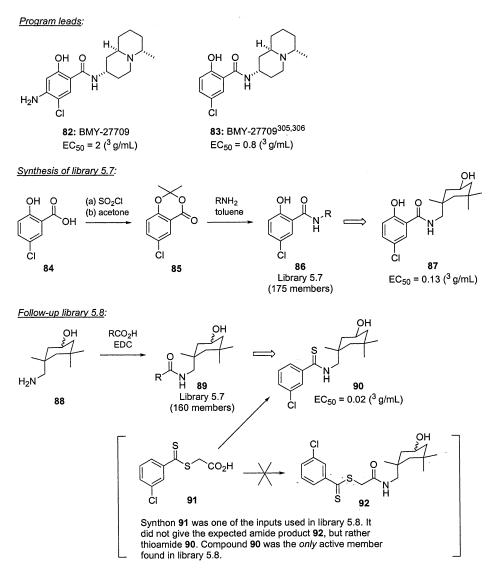
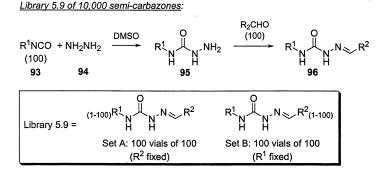


Figure 9. Inhibitors of influenza virus fusion.⁶⁸

The primary amide was next examined (**67B**), and it was established that this functionality could be removed, yielding **68** ($pK_i = 7.7$). This compound displayed a modest loss in antagonist activity and a substantial decrease of plasma clearance, doubling the intravenous half-life. Further modifications to this region of the molecule via **67C**,**D** and shortening the length of the 3-(5-oxohexanoamide) substituent **67E** gave at best modest reductions in potency without any improvement in pharmacokinetic parameters. Attempted optimization of **66** highlights difficulties experienced in simplifying complex, high molecular weight leads and underscores preferred library designs that yield simpler low molecular structures as starting points for optimization.

CNS Library Design: Glycine Transporter-2 (GlyT2) Inhibitors.^{20,49} The design and synthesis of a 3042-member library (library 4.8) on REM resin was carried out at Organon Laboratories (Figure 7).²⁰ Designated as a general-purpose library for screening against diverse molecular targets, its physiochemical properties were biased toward central nervous system (CNS) permeability. This was accomplished by incorporating functionality and molecular properties found in CNS-active drugs. An analysis of the frequency of structural fragments present in 373 orally bioavailable CNS agents revealed that aromatics and amines occur respectively in 83% and 63% of the drugs, followed by alcohols/phenols (21%), esters/aldehydes (\sim 10%), and acids (4%). In addition, the molecular property distribution of these marketed CNS-active drugs are shown in Table 13. Considering these statistical data, library 4.8 was designed to contain a diverse collection of arylamines whose molecular property distribution fell within the above-stated property limits.

Approximately 20 g of 13 phenol amine scaffolds, 4 commercial amines (**70** and **71**), and 9 custom amines (structures not disclosed) were loaded onto REM resin to give ca. 12 g of each resin-bound scaffold **72**. The pheno-lamine cores served to ensure that each library member contained at least one aryl ring and one basic nitrogen atom and provided an avenue for bidirectional derivatization. Chemistry was carried out in either IRORI MicroKans or in Syro II reactors. To maximize chemical diversity within the library, five phenolic OH derivatization chemistries (Mitsunobu, Suzuki, acylation, sulfonylation, and carbamoylation) were employed in combination with three different cleavage strategies ((i) alkylation–Hofmann elimination, (ii) AlCl₃-



Screening library 5.9 reveals by-product 98 as the active constituent:

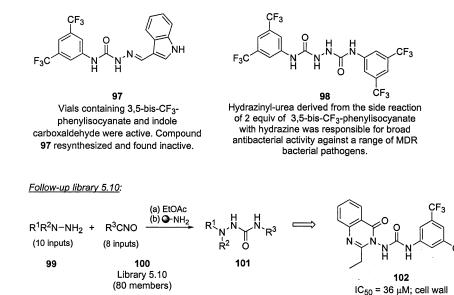


Figure 10. Hydrazinyl-urea byproduct identified as potent antibacterial agent.²⁶⁸

promoted aminolysis, and (iii) base-mediated esterolysis). The library chemistry is the most sophisticated reported to date on REM resin. This approach provided over 3000 tertiary amines containing phenol, aryl ether, aryl ester, aryl sulfonate, biaryl, and propionyl ester and amide functionality. The molecular property distribution of the final library (Table 13) indicated that the majority of compounds are predicted to penetrate into the CNS. Finally, screening library 4.8 against the human GlyT-2 afforded arylamine **78** (IC₅₀ = 1.0 μ M). The physiochemical properties of **78** are largely consistent with a CNS-drug-like profile, although they are at the upper end of the desired range and exceed the preferred number of H-bond acceptors and rotatable bonds. No CNS bioavailability data were reported for **78**.

In a separate screening effort, the Organon group discovered amide **79** (IC₅₀ = 214 nM) to inhibit [³H]-glycine uptake into CHO cells stably expressing the human GlyT-2 protein (GlyT-2 assay).⁴⁹ GlyT-2 is one of two major glycine transporters localized to the spinal cord and brain stem. Glycine is an inhibitory neurotransmitter producing muscle relaxation. Selective inhibition of the transporter is hypothesized to lead to an increase in synaptic levels of endogenous glycine and therefore represents an opportunity for developing a novel muscle relaxant, anesthetic, and/or analgesic. A

"hit-to-lead" program was initiated to increase the potency of 79 while maintaining transporter/receptor selectivity and a CNS-like drug profile. Rapid SAR exploration was achieved by generating library 4.10 (80) via solution-phase methodology. A total of 16 commercially available and custom diamines were derivatized with 18 inputs including 16 carboxylic acid chlorides, 1 sulfonyl chloride, and 1 isocyanate. The two-dimensional array of inputs allowed simultaneous investigation of the structure-activity requirements for cyclic versus acyclic diamines, diamine ring size, α - and β -substitution in the ethylenediamine unit, amide versus another type of linkage, and aromatic ring substitution. Chemistry was performed in four 96-well plates, yielding 10-15 mg quantities of each library member (288 compounds) and purities in excess of 80% (LC/MS). One of the more interesting analogues obtained from screening the library in the GlyT-2 assay was amide 81. It displayed an $IC_{50} = 16 \text{ nM}$ against GlyT-2, >1000-fold selectivity versus GlyT-1 and a panel of receptors, and physiochemical properties consistent with known CNS agents.

biosynthesis inhibition, S. aureus

Influenza Virus Fusion Inhibitors.⁶⁸ Salicylic amide **82** (EC₅₀ = 2^{3} g/mL) was a recently identified as an inhibitor of the H1 and H2 subtypes of influenza A virus strains in cell culture (Figure 9).^{305,306} The biological activity is thought

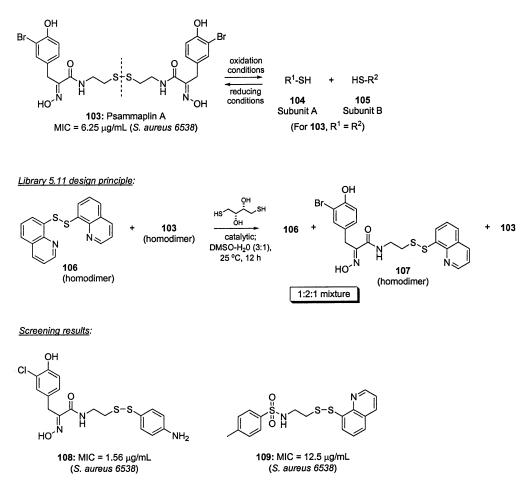


Figure 11. Disulfide library 5.11 based on psammaplin A.^{180,181}

to interfere with virus infectivity by preventing the low-pHinduced conformational rearrangement of hemagglutinin into its fusogenic state and hence blocking obligate virus and host cell fusion. Early SAR work established the importance of the salicylic hydroxyl group, the nonessential anilino group, and stereochemistry of the quinolizidine ring as embodied in analogue 83 (EC₅₀ = 0.8^{3} g/mL). In an effort to increase the potency of the series, to broaden the activity to include H3 influenza subtypes, and to simplify the structure, a twopart sequential optimization program was undertaken. First, surrogates for the quinolizidine ring were sought. Library 5.7 (86; 175 members) was prepared in solution by reacting isopropylidine 85 with a range of primary amines (3-fold excess). This led to amide 87 (EC₅₀ = 0.13^3 g/mL), a substituted cyclohexylmethylamine no longer containing a basic nitrogen. Second, library 5.8 (89), designed to explore the SAR of the salicylic acid, was constructed using the optimized cyclohexylmethylamine synthon from 86. Some 160 amides were synthesized, meeting an 80% purity cutoff for screening. This second library produced only a single active member derived from carboxylic acid 91. The expected product from the reaction was amide 92. However, NMR and MS analysis confirmed the formation of thioamide 90, its formation occurring via an alternative coupling mode between the amine and dithioester carbonyl. Thioamide 90 was a potent inhibitor of influenza viral fusion (EC₅₀ = 0.02^{-3} g/mL), and its unanticipated formation opened up a new window for future SAR development.

Urea-Based Antibacterials.²⁶⁸ A second example of the isolation and identification of a biologically active byproduct from a library was reported by Wilson (Figure 10).²⁶⁸ A 10 000 member mixture-library of semicarbazones (96; library 5.9) was synthesized in solution as shown in Figure 10. Hydrazine 94 was acylated with isocyanates 93 to form semicarbazides 95, which were then condensed with aldehydes to yield semicarbazones 96. Indexed library 5.9 was prepared in two sets of 100 vials each. The first set (A) contained 100 isocyanate inputs per vial and the second set (B) 100 aldehyde inputs. The vials were diluted with DMSO to a compound concentration range of 100-500 mg/mL. Library 5.9 was screened in cell wall biosynthesis and bacterial growth inhibition assays. Two particular vial sets demonstrated dual activity: set A defined by 3,5-bis-(trifluoromethyl)phenyl isocyanate; set B defined by indole-3-carboxaldehyde. This suggested that the activity was due to the cross-product, semicarbazone 97. However, upon resynthesis, its activity did not confirm it. Careful analysis of the active vial from set A revealed two minor byproducts corresponding to the 3,5-bis(trifluoromethyl)phenyl semicarbide (95) and urea 98, where hydrazine had reacted with 2 equiv of isocyanate. Synthesis of these two compounds confirmed biological activity exclusively due to 98. A followup library 5.10 was synthesized to develop its SAR. Urea 102 was one of the more potent compounds obtained from this library.

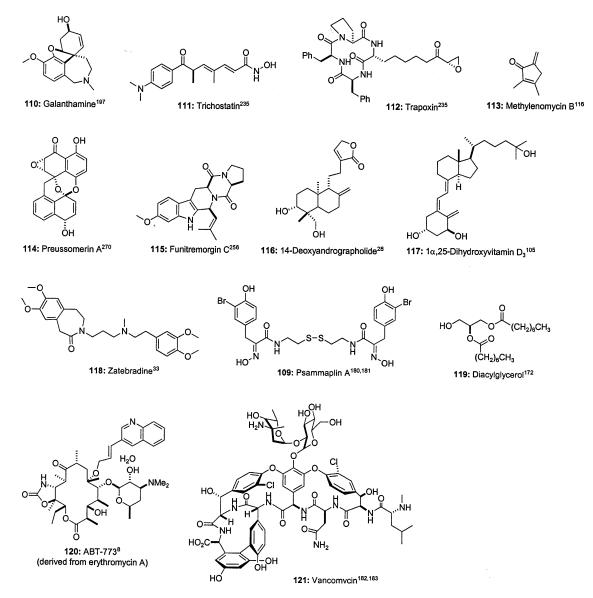


Figure 12. Natural products as library templates in 2001.

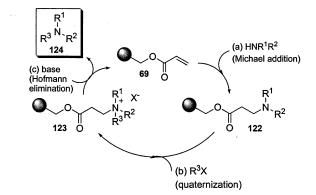
Psammaplin A Based Antibacterials (Natural Product Templates).^{180,181} Psammaplin A 103 is a symmetrical bromotyrosine-derived disulfide natural product exhibiting in vitro antibacterial activity against methicillin-resistant S. aureus (MRSA). In an effort to gather SAR information and gain insight into its mechanism of action, the 3828-member library 5.11 of homodimeric and heterodimeric disulfides was prepared (Figure 11). A novel combinatorial disulfide exchange strategy was devised in which a 1:1 mixture of two homodimeric disulfides (e.g., 106 and 103) was scrambled in the presence of a catalytic amount of dithiotreitol (0.15 equiv, DMSO/H₂O (3:1), 25 °C, 12 h). Under thermodynamic control, the heterodimeric disulfide (e.g., 107) was generated as part of a 1:2:1 mixture of products (106/107/103). Eightyeight custom and commercially available homodimeric disulfides were scrambled in a two-dimensional 96-well plate format. LC/MS analysis of ca. 100 wells showed the formation of expected mixtures of products in excellent purity. Screening the library directly without purification of products against various strains of MRSA afforded several active wells. Resynthesis of these screening hits confirmed

a number of heterodimeric disulfides, in some instances with increased antibacterial potency over the natural product (e.g., **108** and **109**). Further optimization of the leads was reported in a subsequent report.¹⁸¹ It was established that the mechanism of action of **103** and related analogues is not attributed to the inhibition of DNA gyrase as previously proposed. Circumstantial evidence was presented suggesting that **103** does interact with a specific, albeit presently unknown, molecular target(s) as opposed to acting through a nonspecific redox-type mechanism.

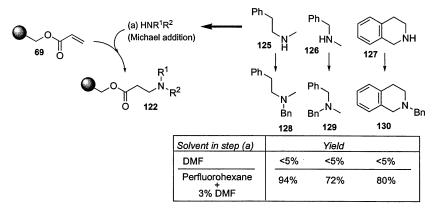
Other natural products serving as inspiration for library synthesis in 2001 are shown in Figure 12.

2. Solid- and Solution-Phase Transformations: Libraries without Disclosed Biological Activity (Tables 6–10)

Advances in REM Resin Methodology.¹⁶⁹ As exemplified in the preparation of the CNS-biased library 4.8, REM resin methodology is an efficient protocol for the solid-phase synthesis of tertiary amines (Figure 13A). REM chemistry was first introduced by Brown, Rees, Rankovic, and Morphy



B. Fluorocarbon solvents facilitate Michael addition on REM resin:169



C. Two-resin method for cleavage of amines from REM resin:10

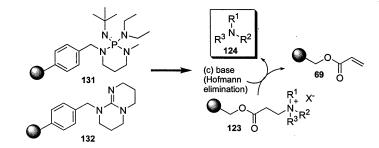


Figure 13. Advances in REM resin methodology.^{10,169}

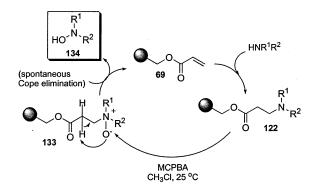


Figure 14. Substituted hydroxylamines from REM resin.²¹⁸

at Organon in 1997.³⁰⁷ It proceeds via the Michael addition of amines to resin-bound acrylate (69-122), formation of quaternary ammonium salt 123 with an alkyl halide, and then upon exposure to mild base, Hofmann elimination liberating tertiary amine 124 and regeneration of resin 69. Continued

research at Organon led to the observation that perfluorous organic solvents allow for a large reduction in both reaction time and the amounts of amine required for the Michael reaction (Figure 13B). Dramatic increases in product yields were also realized. Given the immiscibility of perfluorocarbons with common organic solvents, the observed rate and yield enhancements are presumably due to trapping amine reagents inside the bead, a so-called "reagent concentration effect". This concept of "fluorcarbon accelerated supported transformations" (FAST) may have broad application in solid-phase synthesis. In the present REM resin study, three amines 125-127 were reacted with 69 in a variety of solvents and combinations thereof: DMF, DMSO, neat amine, water, hydrocarbons, and perfluoroalkyls and perfloroaromatics. Perfluoroaromatic solvents did not give any product after resin quaternization with benzyl bromide and Hofmann elimination (128-130). Reaction of 69 with 4 equiv (versus 20 equiv) of amine for 2 h was sufficient to effect Michael

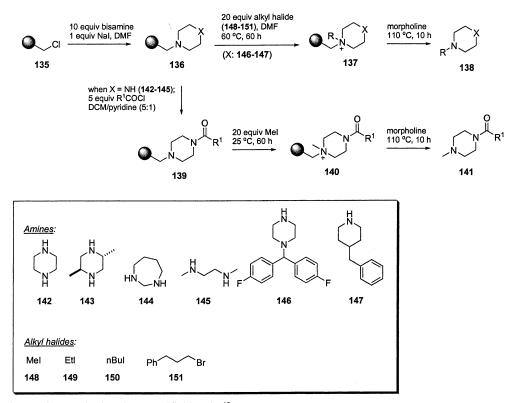
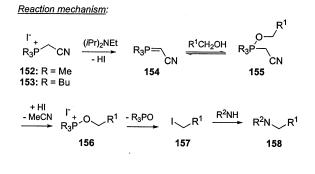


Figure 15. Traceless amine synthesis using Merrifield resin.45



Examples of alkylation products (* = resin attachment point):

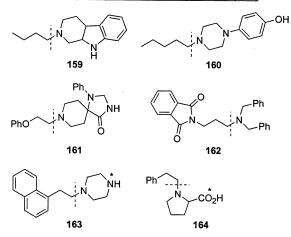


Figure 16. Alkylation of secondary amines with primary alcohols using (cyanomethyl)trialkylphosphonium iodides.²⁸⁸

addition. The perfluorohexane containing 3% DMF appeared to be the optimal solvent combination. Perflurous solvents also had a positive effect on the quaternization and Hofmann elimination steps; details are forthcoming.

Alhambra reported that resin-bound phosphazene **131** or quanidine **132** significantly improved the yield of the Hofmann elimination step (Figure 13C).¹⁰ Under standard cleavage conditions with Et_3N or $(iPr)_2EtN$, the tertiary amine products may be contaminated with Et_3N or $(iPr)_2EtN$ hydrobromide or hydroiodide salts. Postcleavage extraction of SPE chromatography can be used to purify products, but this is labor-intensive with large sets of compounds. The advantage of a two-resin cleavage method is that it avoids salt formation and gives products of superior purity. Resinbound base **131** and in particular **132** were reported to be superior to previously employed resin-bound diisopropylbenzylamine.

Substituted hydroxylamines **134** were prepared using REM resin (Figure 14).²¹⁸ After Michael addition, resin **122** is treated with *m*-chloroperbenzoic acid (MCPBA) at 25 °C to furnish the corresponding amine oxide **133**. Resin **133** undergoes spontaneous Cope elimination and release of hydroxylamine.

New Methods for Tertiary Amine Synthesis.^{45,96,288} Merrifield resin was utilized in a traceless synthesis of tertiary amines (Figure 15). Substitution of the benzylic chloride in 135 with secondary amines (X not equal to NH) using NaI as a catalyst afforded the corresponding resin-bound amine 136. Quaternization employing MeI took place at 25 °C in DMF, while use of alkyl halides 149–151 required heating at 60 °C for up to 60 h. Dequaternization and release of the tertiary amine was effected by heating resin 137 with morpholine at 110 °C for 10 h. The selection of morpholine as the preferred cleavage reagent is due to its superior resinswelling property. The utility of the chemistry was further demonstrated by constructing a library of 16 compounds in IRORI MacroKans in which secondary diamines (142–145) A. Alkyl halide formation:200

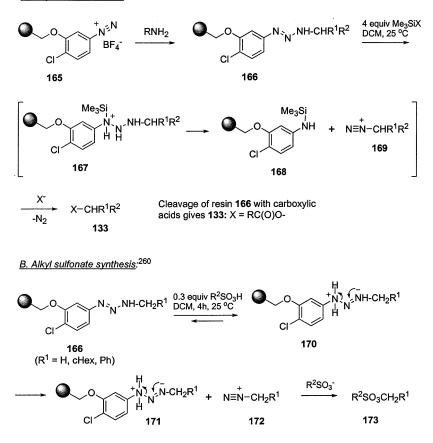


Figure 17. Application of resin-bound triazene in solid-phase synthesis.

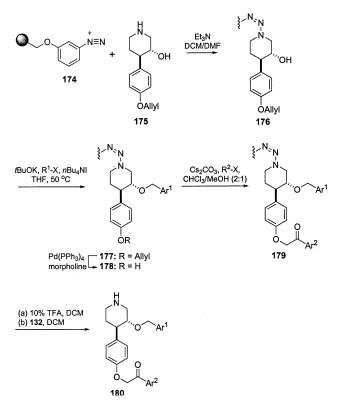


Figure 18. Synthesis of 3,4-disubstituted piperidines using triazene linker.⁴⁴

attached to resin were N-acylated, converted to their methylammonium salts 140, and cleaved with morpholine to give methyl tertiary amines 141. Yields averaged \sim 75% with product purities in excess of 90%.

Zaragoza and Stephensen at Novo Nordisk succeeded in utilizing (cyanomethyl)trialkylphosphonium iodides as efficient reagents for alkylation of amines with alcohols in solution and on solid phase (Figure 16).²⁸⁸ This operationally simple method is carried out by adding (cyanomethyl)trimethylphosphonium iodide 152 (1.2 equiv) to a mixture of a primary alcohol (1.05 equiv), a secondary amine (1 equiv), and DIEA (1.3 equiv) in propionitrile. The reaction mixture was heated to 90 °C for 2 h. An aqueous solution of K₂CO₃ was then added to quench the reaction, and the tertiary amine 158 was extracted with a suitable solvent (EtOAc) and purified by standard techniques. The mechanism of the reaction involves deprotonation of the phosphonium salt 152 (153) and reaction with alcohol to yield species 155. Thermolysis of 155 in the presence of DIEA hydroiodide leads to P-C bond cleavage, release of acetonitrile, and formation of alkoxyphosphonium salt 156. Decomposition of 156 yields primary iodide 157, the amine alkylating agent. In a solid-phase variant, resin-bound proline (Wang resin) was treated with phosphonium salt 153 (6 equiv) and a solution of alcohol (7 equiv) and DIEA (8 equiv) in acetonitrile. The reaction mixture was shaken at 80 °C for 24 h, filtered, extensively washed, and then cleaved with TFA/DCM to furnish the tertiary amine in good yield $(\sim 75\%)$ and high purity. Examples of amines 159–164 produced using this protocol (in solution and on solid phase)

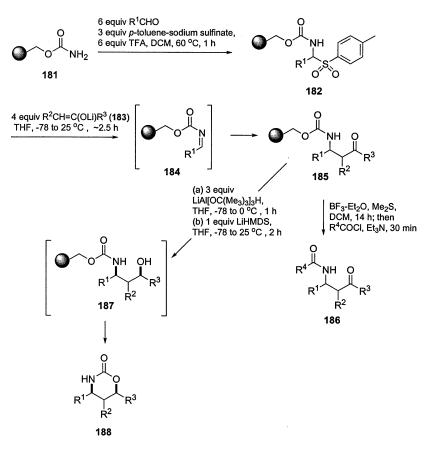


Figure 19. Resin-bound acylimines.²²

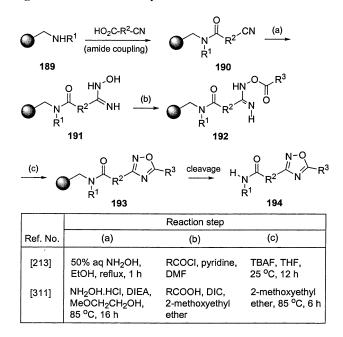


Figure 20. 1,2,4-Oxadiazole synthesis on solid support (from resinbound nitriles).

are illustrated in Figure 16 (amine and alcohol retrocoupling indicated at dashed line).

Andersson and co-workers described a novel solid-phase synthesis of tertiary methylamines involving iodide- or SmI₂based cleavage of the N–O bond of resin-bound alkoxyammonium salts.⁹⁶ These were prepared via the borane reduction of resin-bound oximes to the corresponding *O*-polystyrene-

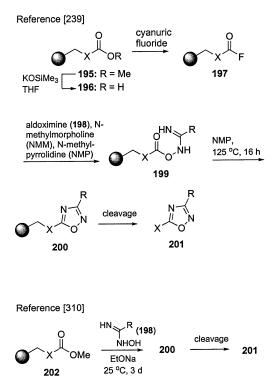


Figure 21. 1,2,4-Oxadiazole synthesis on solid support (from resinbound esters).

linked hydroxyamine followed by N-alkylation and quaternization with MeOTf. Purity of the exemplified amine was >90%. The reaction reportedly has broad scope, but details were not disclosed.

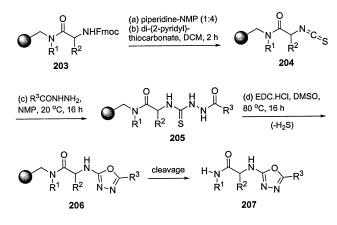


Figure 22. 1,3,4-Oxadiazole synthesis on solid support.¹²⁵

Resin-Bound Triazenes as Reagent and Linker.^{44,200,260} Brase and co-workers developed a method for attaching secondary amines onto and detaching secondary amines from solid-support via T2 triazene linkages, e.g., 166.313 During investigations of amide bond formation on the resin, it was observed that in cases where acylation was problematic, subsequent cleavage of material from resin with trimethylsilyl chloride (TMS-Cl) gave alkyl chlorides. This observation led to the development of a solid-phase synthesis of alkyl halides and esters from primary amines. The mechanism of the reaction is presented in Figure 17A. Reaction of stable diazonium resin 165 with a primary amine proceeded smoothly to give disubstituted triazenes 166. Treatment of 166 with a 10% solution of TMS-Cl in DCM for a few minutes at 25 °C, filtration, and evaporation of solvent afforded the alkyl chlorides in excellent yield and purity. TMS-Br and TMS-I gave the corresponding alkyl halides. Alkyl acetates and trifluoroacetates are produced from nucleophilic cleavage with acetic and trifluoroacetic acids, respectively. The method can also be used for the mild and selective synthesis of sulfonic acid esters 173 from sulfonic acids or sodium sulfonates (Figure 17B).²⁶⁰ Twelve examples of sulfonate esters were reported with yields ranging from 66% to 91%.

One noteworthy application of the triazene linker in library synthesis was reported by Rich in the preparation of putative aspartic acid protease inhibitors (Figure 18).44 Triazene-linked piperidine 176, prepared by reacting diazonium resin 174 with 175, was smoothly alkylated with three different benzyl bromides using potassium tert-butoxide as base in THF at 50 °C. Pd⁰-mediated deallylation of **177** occurred smoothly to give phenols 178. Alkylation of 178 with aryl bromomethyl ketones (five inputs) required optimization. Standard reaction conditions (base, DMF) gave only minimal conversion to 179 even in the presence of nBu_4NI . An obscure patent citation describing K2CO3-mediated O-alkylation of phenols in a 2:1 mixture of CH₃Cl/MeOH³¹⁴ prompted the use of these conditions in the current synthesis and afforded conversions of $\sim 90\%$ to the desired bis-ether 179. Library members 180 were generated upon exposure of 179 to 10% TFA/DCM and free-basing of the trifluoroacetate salts with resin-bound guanidine 132.

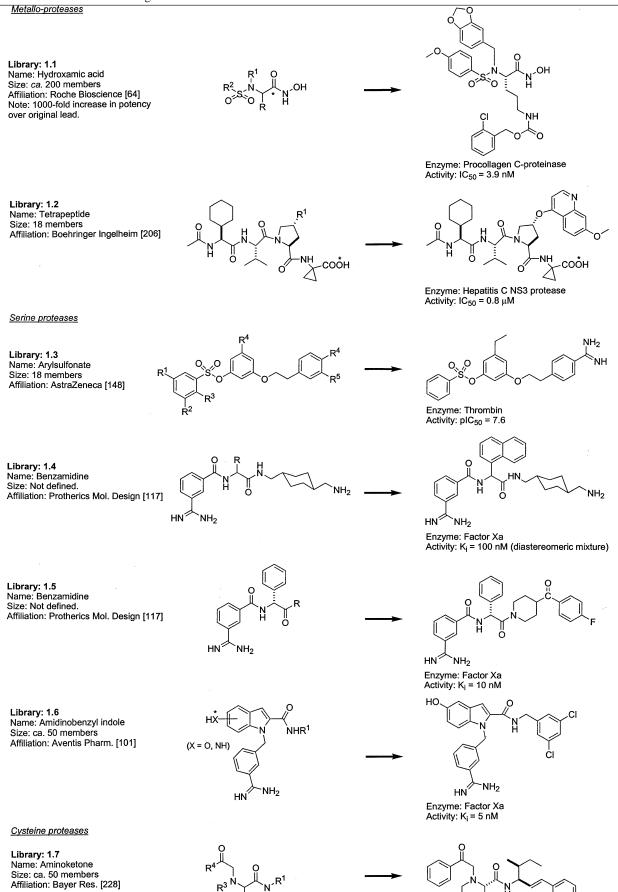
Resin-Bound *N***-Acylimines.**²²² Acylimines, the neutral congener of the *N*-acyliminium ion, are generated in situ from

 α -amino sulfones. Acylimines are reactive intermediates known to undergo Mannich-type addition with a wide variety of nucleophiles including ketone enolates, Reformatsky reagents, nitromethane anion, and vinyl and akynyl organometallic reagents. Enders adapted the synthesis of α -amino sulfones (182) to solid phase by condensing carbamate linker **181** with an aldehyde and *p*-toluenesodium sulfinate in the presence of TFA (Figure 19). Upon treatment of with an excess of base, e.g., ester enolate 183, and elimination of lithium p-toluolsulfinate, acylimine species 184 was produced, which immediately underwent a Mannich reaction to give the resin-bound *N*-acyl- β -amino ester **185**. Dimethyl sulfide promoted BF₃/Et₂O cleavage of 185 afforded the corresponding amine, which may optionally be acylated to give 186. Stereoselective reduction of the ketone carbonyl in 185 generates alcohol 187 and yields cyclic carbamates **188** via intracyclative cleavage. The percent diastereomeric excess (% de) for the reduction generally exceeded 94%, and purity of cleaved products was >85%. Treatment of 182 with allylzinc or PhMgCl gave the expected Mannich addition products.

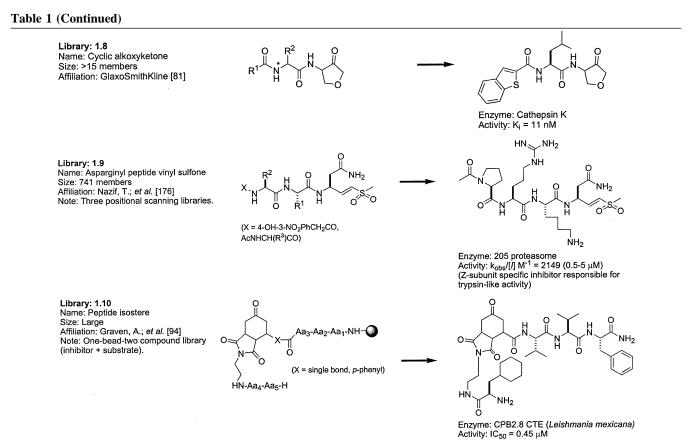
Oxadiazoles.^{85,125,203,213,311} 1,2,4-Oxadiazoles are considered metabolically stable ester bioisosteres and are found in a large number of biologically active compounds of pharmaceutical interest. Several solid-phase syntheses of this heterocycle have been reported via cyclodehydration of O-acylamidoximes as the key ring-forming step (Figures 20) and 21). Resin-bound O-acylamidoximes have been prepared by one of two ways: (i) condensation of hydroxylamine with resin-bound nitriles followed by acylation (Figure 20);^{85,213,311} (ii) reaction of resin-bound activated carboxylic acids^{203,239} or esters³¹⁰ with aldoximes (Figure 21). Thermal cyclodehydration^{239,308,310,311} of *O*-acvlamidoximes has been the preferred option for heterocyclic ring formation, but recently Kenner (solid-phase),²¹³ then Gangloff (solution-phase),85 used tetra-N-butylammonium fluoride (TBAF) to cyclize O-acylamidoximes 192 at room temperture (192 and 193; Figure 20). TBAF-mediated cyclodehydration is compatible with a range of functionality affording diverse collections of 3,5-disubstituted 1,2,4oxadiazoles.

The synthesis of isomeric 1,3,4-oxadiazoles **207** was also reported (Figure 22).¹²⁵ Resin-bound thioisocyanates **204** were converted to acylthiosemicarbazides **205**. A survey of dehydrating reagents and conditions to convert **205** to **206** was carried out, and optimal cyclodehydration was effected using EDC·HCl in DMSO at 80 °C. Yields for **207** (six examples) ranged from 44% to 92%; product purities were in excess of 90%. Previous library-based methods for 1,3,4oxadiazole synthesis include the dehydration of 1,2-diacylhydrazines on solid phase and in solution phase, the latter employing polymer-supported Burgess reagent in combination with microwave heating as the dehydration conditions.^{309,312}

Acknowledgment. Sincere appreciation is expressed to Becca Schaefer for her expert assistance in chemical structure drawing and in the preparation of other portions of this manuscript.



Enzyme: Cathepsin K Activity: K_i = 230 nM



^a The asterisk (*) indicates point of attachment to the resin.

Table 2. Chemical Libraries Targeted for Nonproteolytic Enzymes^a

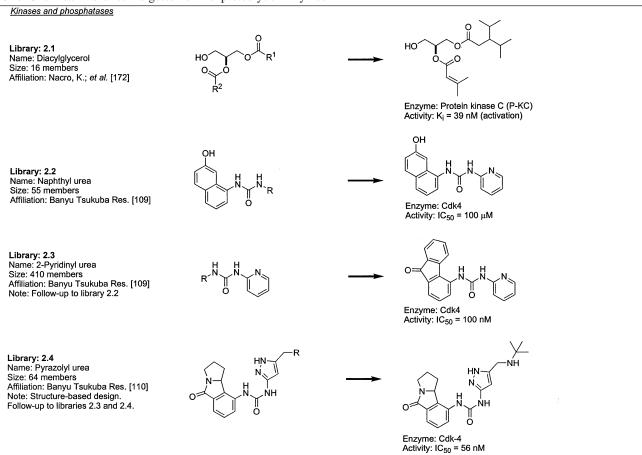


Table 2 (Continued)

Kinases and phosphatases

Library: 2.5 Name: Heterocyclic urea Size: ca. 1000 members Affiliation: Bayer Res. [227]

Ar^{-N}

R²

R²

.OPO3⁻⁻⁻

з́н

ċΝ

H-Aa₄-Aa₃-Aa₂-Aa₁-ÔH

R

Library: 2.6 Name: Sulfonylated aminothiazole Size: 35 members Affiliation: Wipf, P.; *et al.* [269]

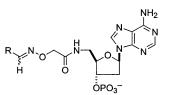
Library: 2.7 Name: Tyrosine phosphate Size: 176 members Affiliation: Shen, K.; *et al.* [225]

Library: 2.8 Name: Diaryl ether Size: Not specified Affiliation: Merck Res. Lab. [154]

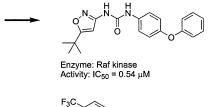
Library: 2.9 Name: Tetrapeptide Size: 331,776 members Affiliation: Henlin, J. M.; *et al.* [103] Note: Mix and split synthesis and deconvolution.

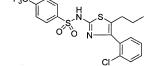
Library: 2.10 Name: Purine Size: 275 members Affiliation: Verdugo, D. E.; *et al.* [258]

Library: 2.11 Name: Nucleotide oxime Size: 447 members Affiliation: Armstrong, J. I.; *et al.* [12]

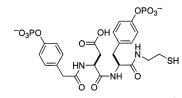




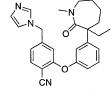




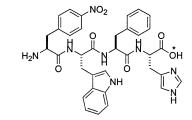
Enzyme: Cdc25B (dual specificity phosphatase) Activity: IC_{50} = 14 μ M



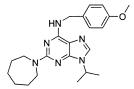
Enzyme: Protein-tyrosine phosphatase 1B Activity: $K_i = 2.4$ nM; selective against a panel of phosphatases



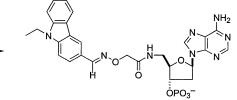
Enzyme: Farnesyl-protein transferase (FPTase) Activity: IC₅₀ = 2.9 nM



Enzyme: S-Farnesyltransferase Activity: $K_i = 2.0 \ \mu M$



Enzyme: Estrogen sulfotransferase Activity: IC₅₀ = 500 nM



Enzyme: Estrogen sulfotransferase Activity: 80% inhibition at 200 μ M

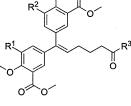


Bacterial and viral enzymes Library: 2.12 соо́н R³HN Name: Trisubstituted pyrrolidine Size: ca. 550 members Affiliation: Abbott Lab. [263] Note: Structure-based design. NR¹R² Library: 2.13 Name: Tetra-substituted pyrrolidine H_2N -OH Size: ca. 20 members Affiliation: Abbott Lab. [263] RHN Note: Follow-up to library 2.12 prepared by solution-phase methodology. Structurebased design. Library: 2.14 Name: N-Hydroxamic acid ЮΗ Size: ca. 20 members Affiliation: Hoffmann-La Roche [11] ö k^2 R3 .R⁴ Library: 2.15 Name: 1,3-Dioxane Size: 7200 members ö 0 О Affiliation: Sternson, S. M.: et al. [235] HO Library: 2.16 Name: Pyrimidone Size: 6 members Affiliation: Botta, M.; et al. [34] НÕ

Library: 2.17 Name: Alkenyldiarylmethane Size: 3 members Affiliation: Xu, G.; et al. [276]

Mammalian enzymes

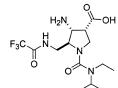
Library: 2.18 Name: Propargylic amine Size: 20 members Affiliation: Conn, C.; *et al.* [57]



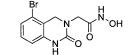
 $R^3 R^4$ NR⁵R⁶ R^1

COOH H₂N

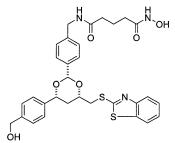
Enzyme: Influenza neuraminidase (A/Tokyo/3/67 virus) Activity: IC₅₀ = 1.3 µM



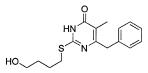
Enzyme: Influenza neuraminidase Activity: IC₅₀ = 0.28 μM



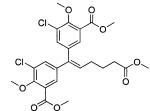
Enzyme: Peptide deformylase (E. Coli) Activity: IC₅₀ = 49 nM



Enzyme: Histone deacetylase Activity: $IC_{50} = 1.2 \ \mu M$



Enzyme: HIV-1 reverse transcriptase Activity: Ki = 70 µM

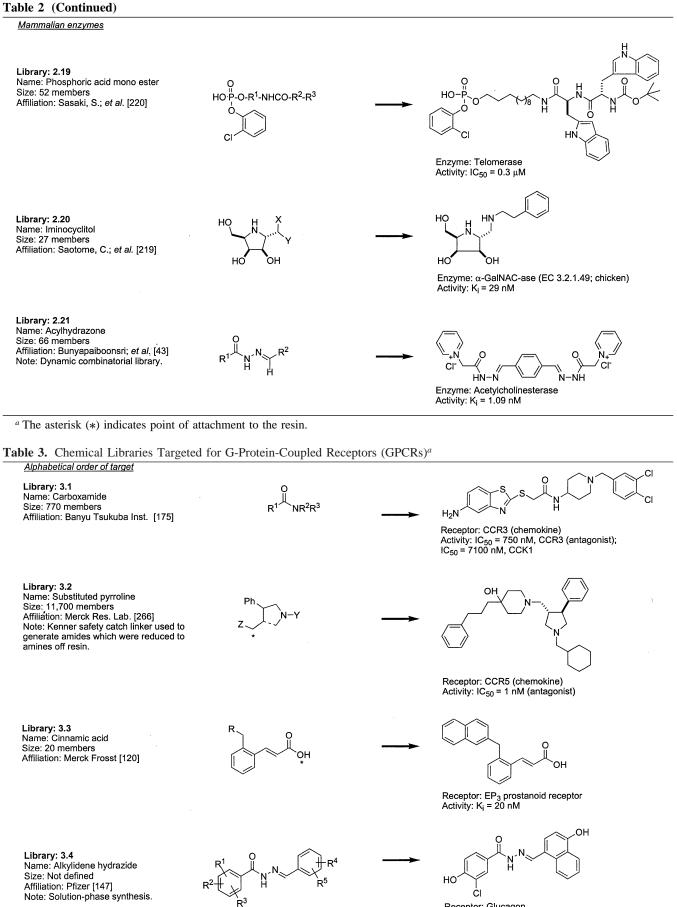


Target: HIV-1 reverse transcriptase Activity: IC₅₀ = 300 nM

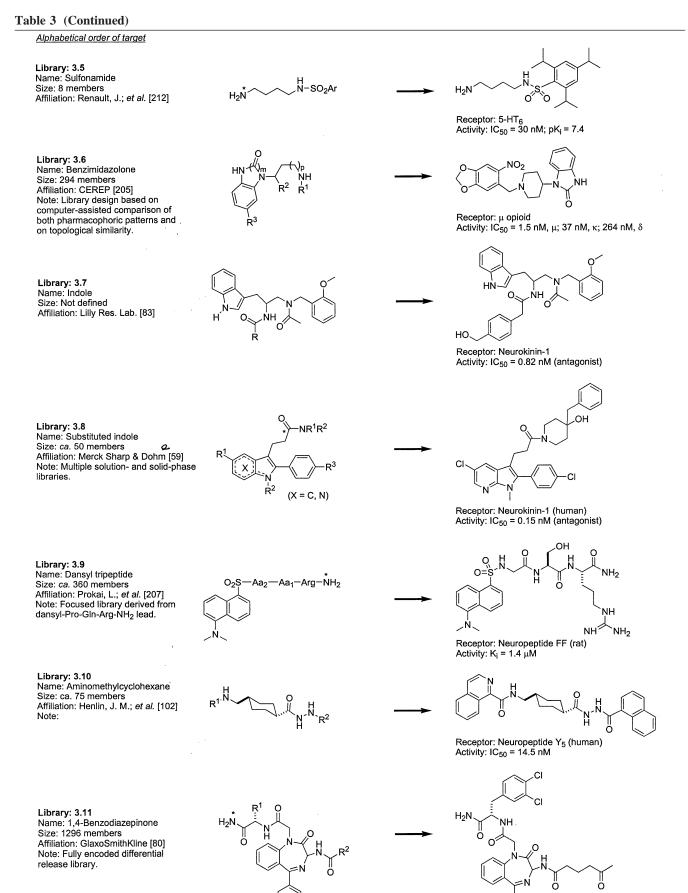
'N C

Enzyme: Amine oxidase (semicarbazide-sensitive) Activity: $K_i = 2.9 \mu M$

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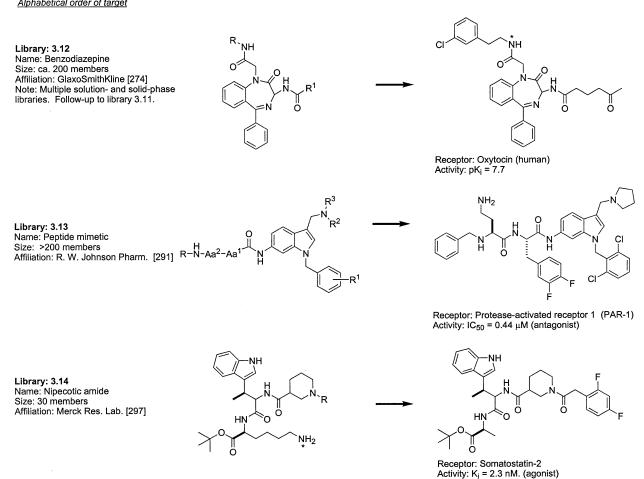
Receptor: Glucagon Activity: $IC_{50} = 0.2 \ \mu M$ (antagonist; inhibition of hyperglycemic effect of glucagon challenge by iv administration in rat)



Receptor: Oxytocin (human) Activity: IC₅₀ = 5 nM (pK_i = 8.1)

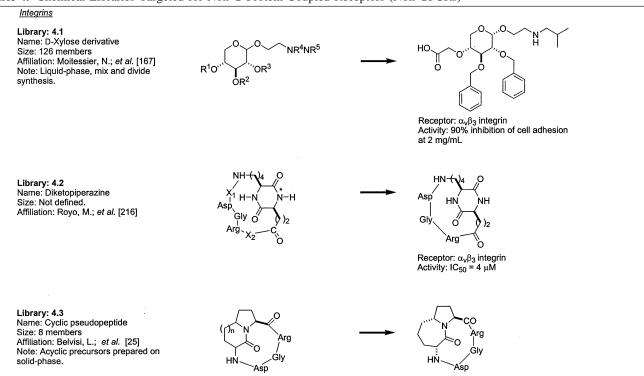
Table 3 (Continued)

Alphabetical order of target



^a The asterisk (*) indicates point of attachment to the resin.

Table 4. Chemical Libraries Targeted for Non-G-Protein-Coupled Receptors (Non-GPCRs)^a



Receptor: $\alpha_{\nu}\beta_3$ integrin Activity: IC₅₀ = 3.7 nM

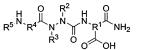
Table 4 (Continued)

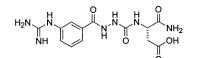


Library: 4.5

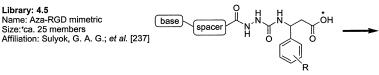
Name: Aza-RGD mimetric Size:'ca. 25 members

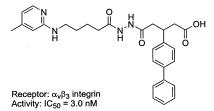
Library: 4.4 Name: Azarurea Size: 990 members Affiliation: Gibson, C.; et al. [88]



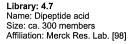


Receptor: $\alpha_{\nu}\beta_3$ integrin Activity: IC₅₀ = 150 nM, $\alpha_{\nu}\beta_3$; IC₅₀ = 7.2 μ M, $\alpha_{\nu}\beta_5$





Library: 4.6 Name: Urea carboxylate Size: ca. 200 members Affiliation: Aventis Pharm. [14]



RSO₂ (or RCO)-Aa₂-Aa₁-CO₂H

 $\overset{R^{3}}{\underset{\scriptstyle \square}{\overset{\scriptstyle N}{\overset{\scriptstyle N}}}}\overset{R^{2}}{\underset{\scriptstyle \square}{\overset{\scriptstyle R}{\overset{\scriptstyle N}{\overset{\scriptstyle }}}}}\overset{R^{1}}{\underset{\scriptstyle \square}{\overset{\scriptstyle \square}{\overset{\scriptstyle \square}{\overset{\scriptstyle N}{\overset{\scriptstyle }}}}}}\overset{O}{\underset{\scriptstyle \square}{\overset{\scriptstyle +}{\overset{\scriptstyle }}}$

HN OH

Receptor: VLA-4 ($\alpha_4\beta_1$ integrin) Activity: IC₅₀ = 1.7 μ M (inhibition of VLA-4 binding to VCAM-1); IC₅₀ = 0.1 μ M (inhibition of VLA-4 binding to fibronectin)

ΩН έo₂ Ö

Receptor: VLA-4 ($\alpha_4\beta_1$ integrin) Activity: IC₅₀ = 7 nM

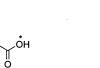
Transporters

Library: 4.8 Name: Aminophenol Size: 3042 members Affiliation: Organon Res. [20] Note: REM resin.

Library: 4.9 Name: Glycine amides Size: Not defined Affiliation: Organon Res. [38] Note: Amine selection based on known serotonin and monoamine uptake inhibitors. Most active inhibitor identified from the library was resolved.

Library: 4.10 Name: Benzamide Size: 288 members Affiliation: Caulfield, W. L.; et al. [49]

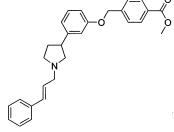
R он spacer N н



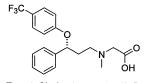
R¹

R¹_N

 k^{2}



Target: Glycine T₂ transporter (human) Activity: $IC_{50} = 1.0 \ \mu M$



Target: Glycine transporter-1b (human) Activity: pIC₅₀ = 6.9

Target: Glycine T2 transporter Activity: IC₅₀ = 16 nM

Reviews

Table 4 (Continued)

lon channels

Library: 4.11 Name: Acylguanidine Size: ca. 200 members Affiliation: CeNeS Pharm.[190]

Library: 4.12 Name: Benzamide Size: Not defined Affiliation: Bristol-Meyers Squibb [150]

Library: 4.13 Name: Zatebradine analog Size: 21 members Affiliation: Organon Lab. [33]

Miscellaneous

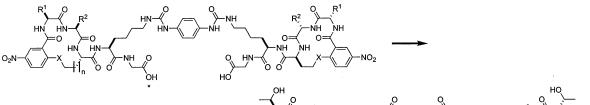
Library: 4.14 Name: N-alkylated thiamine Size: 42320 members Affiliation: Tai, K.-K.; et al. [242] Note: Series of positional scanning libraries.

Library: 4.15 Name: Substituted aryl acid Size: 480 members Affiliation: GlaxoSmithKline [149]

Library: 4.16 Name: Thioisobutyric acid Size: ca. 160 members Affiliation: GlaxoSmithKline [39]

Library: 4.17 Name: Dimeric β-turn peptidomimetic

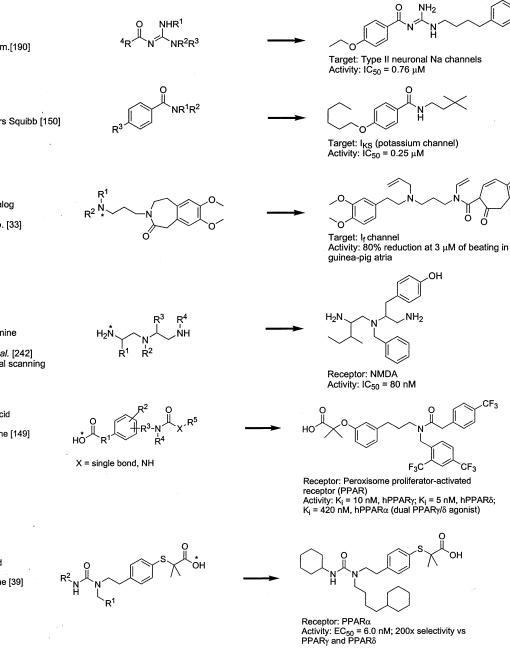
Size: 12 members Affiliation: Zhang, A. J.; *et al.* [289]

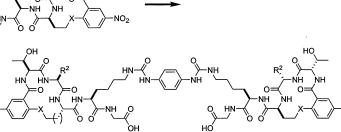


O₂N

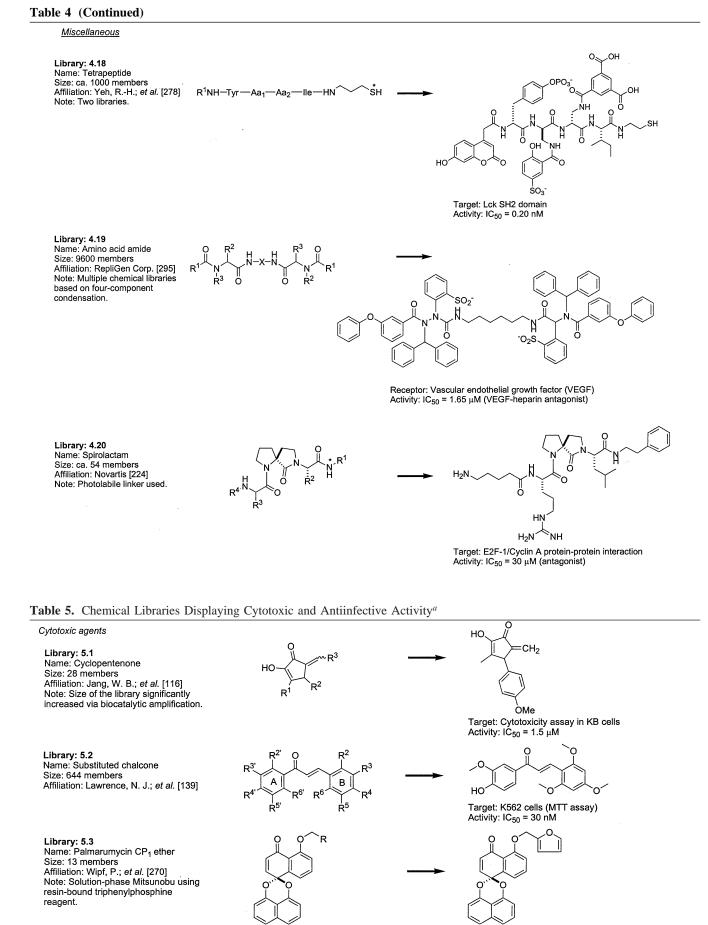
0.

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Receptor: Tyrosine kinase-C for neurotrophin-3 Activity: Increase in extracellular acidification rate over baseline.



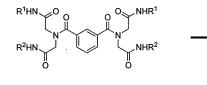
Target: MCF-7 cancer cell line $IC_{50} = 1.1 \ \mu M$

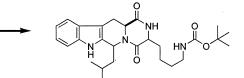


Cytotoxic agents

Library: 5.4 Name: Fumitremorgin analog Size: 42 members Affiliation: van Loevezijn, A.; *et al.* [256]

Library: 5.5 Name: Benzene-1,3-dicarboxylic acid amide Size: 600 members Affiliation: Boger, D. L.; *et al.* [31]

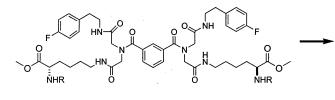




Target: Mitoxantrone accumulation assay (T* human cell line) Activity: Comparable to fumitremorgin

antagonist activity

Library: 5.6 Name: Symmetrical tetraamide Size: 77 members Affiliation: Boger, D. L.; *et al.* [31] Note: Follow-up to library 5.5.



Activity: $IC_{50} = 6 \ \mu M$ (disruption of MMP2- $\alpha v \beta 3$ integrin protein-protein interaction

ĊF3

Antiinfective agents

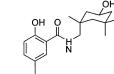
Library: 5.7 Name: Salicylamide Size: 175 members Affiliation: Bristol-Meyers Squibb [68]

Library: 5.8 Name: Cyclohexylmethylamide Size: 160 members Affiliațion: Bristol-Meyers Squibb [68] Note: Follow-up to library 5.7.

Library: 5.9 Name: Hydrazinyl-urea Size: 80 members Affiliation: Procter & Gamble Pharm. [268] Note: By-product from mixture-based synthesis was the active component.







ĊF₂

Target: Influenza virus fusion Activity: $EC_{50} = 0.08$ ³g/mL

ĊΙ

Target: Influenza virus fusion Activity: $EC_{50} = 0.02 \ {}^{3}g/mL$

 $\mathbb{R}^{1} \stackrel{\mathsf{H}}{\longrightarrow} \mathbb{N}^{1} \stackrel{\mathsf{N}_{\mathbb{N}}}{\longrightarrow} \mathbb{N}^{2}$

CF₃ F₃C O ĊF₃

Target: Peptidoglycan biosynthesis (S. aureus) Activity: IC₅₀ = 17 μ M

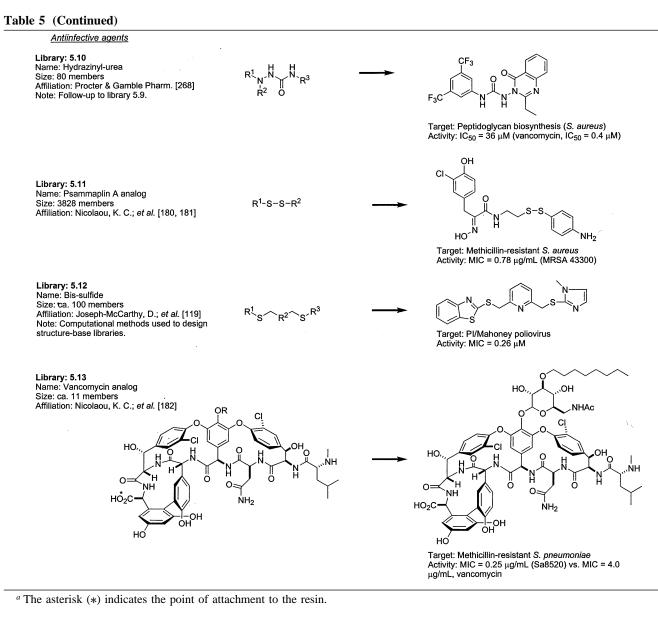
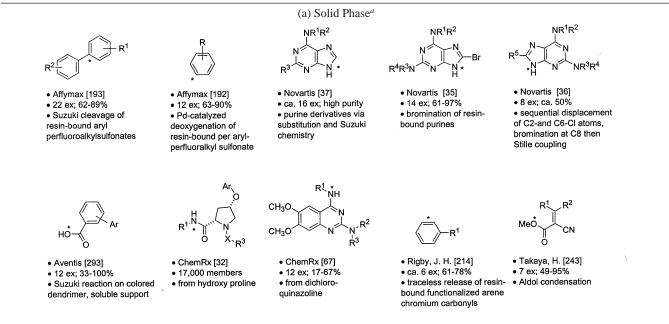


Table 6. Scaffold Derivatization



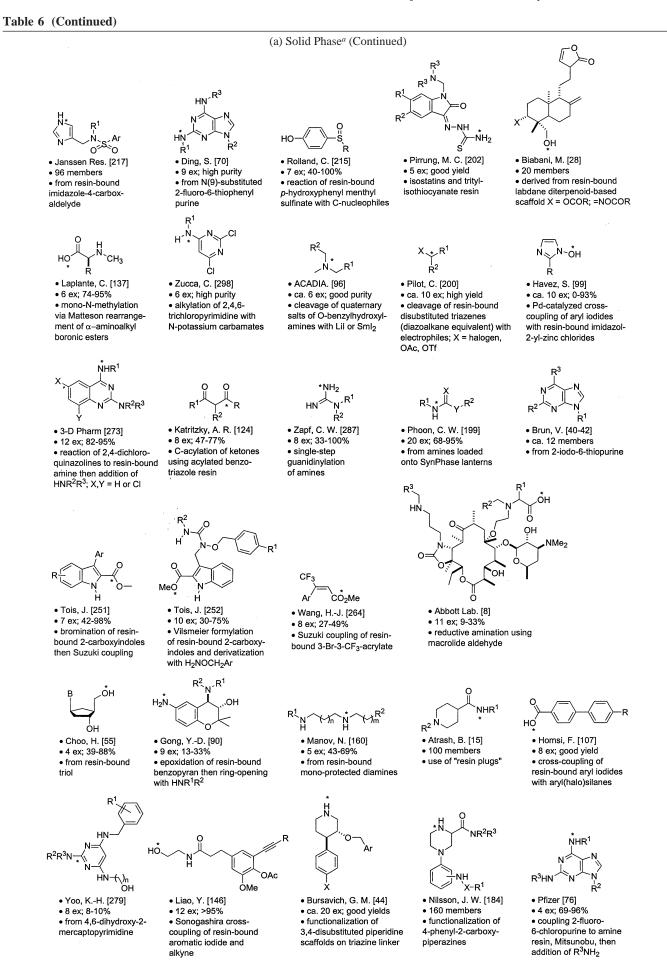
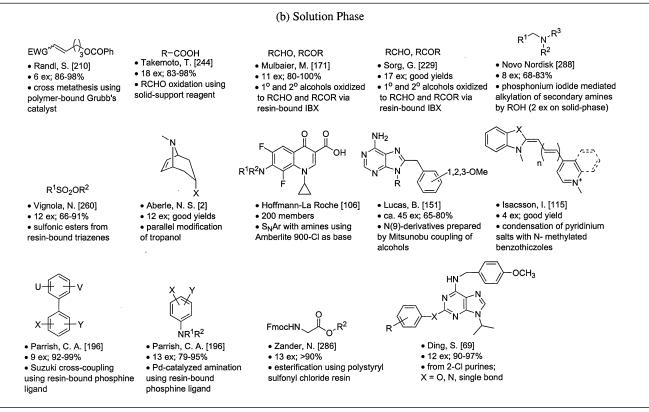
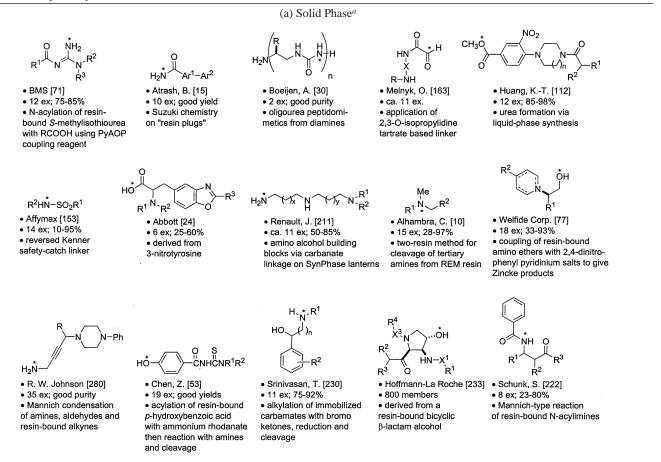


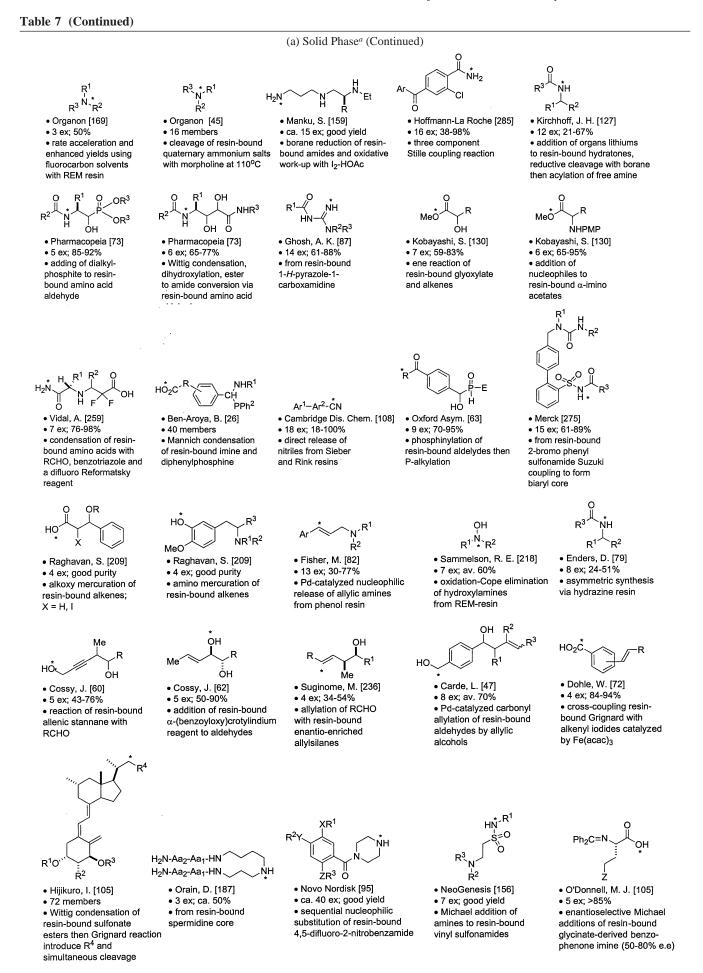
Table 6 (Continued)



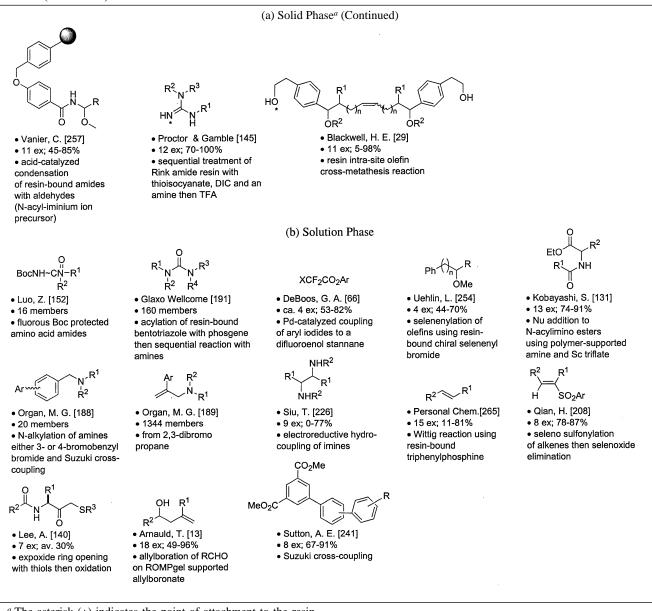
^a The asterisk (*) indicates the point of attachment to the resin.





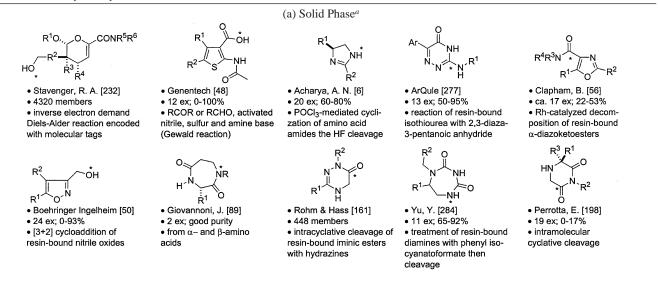


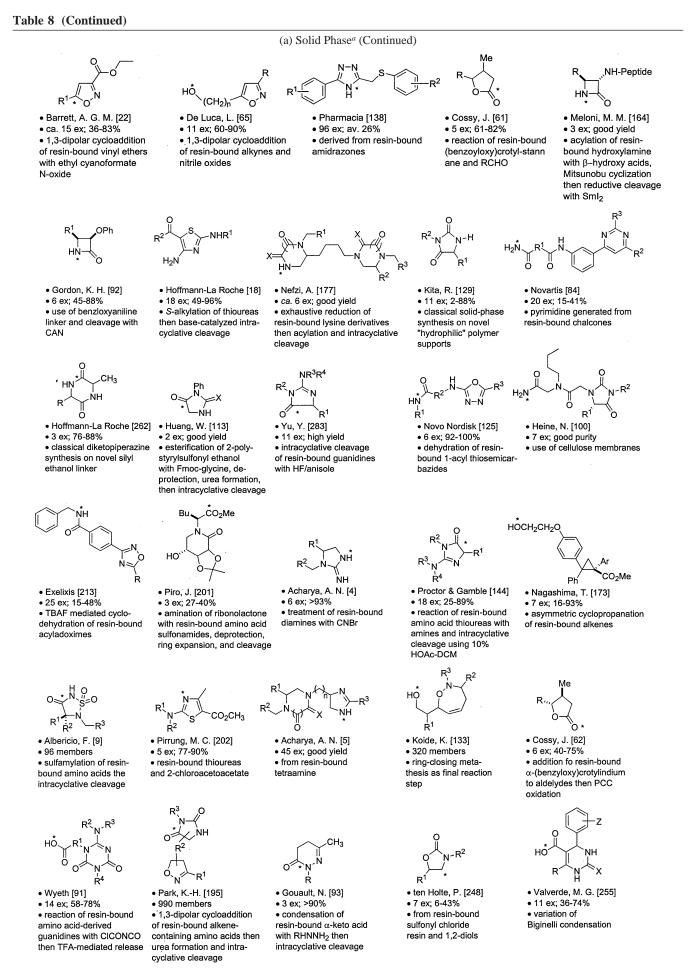


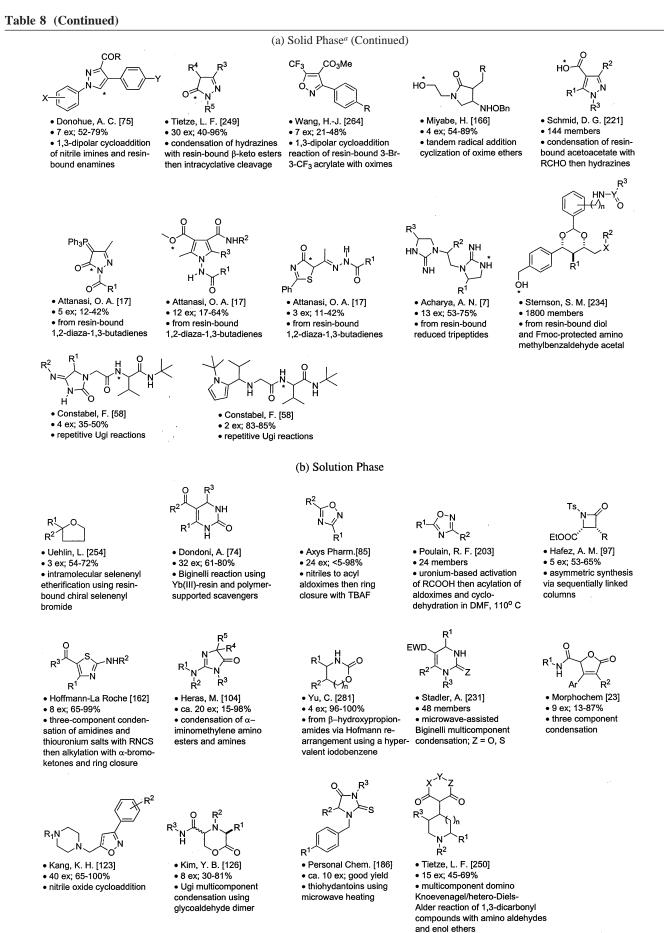


 a The asterisk (*) indicates the point of attachment to the resin.

Table 8. Monocyclic Synthesis





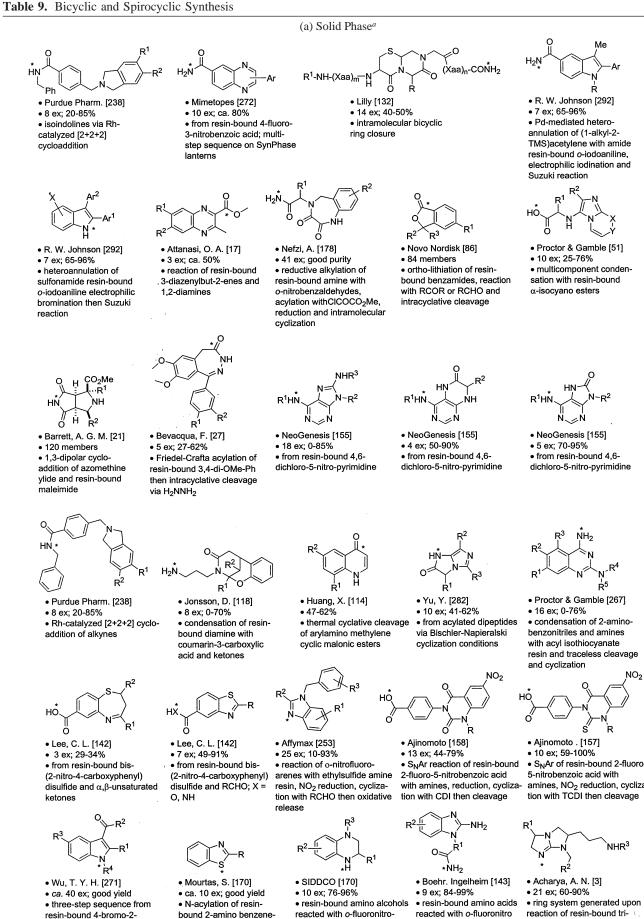


^a The asterisk (*) indicates the point of attachment to the resin.

iodoaniline

thiol then intracyclative

cleavage



reduction, cyclization

then cleavage

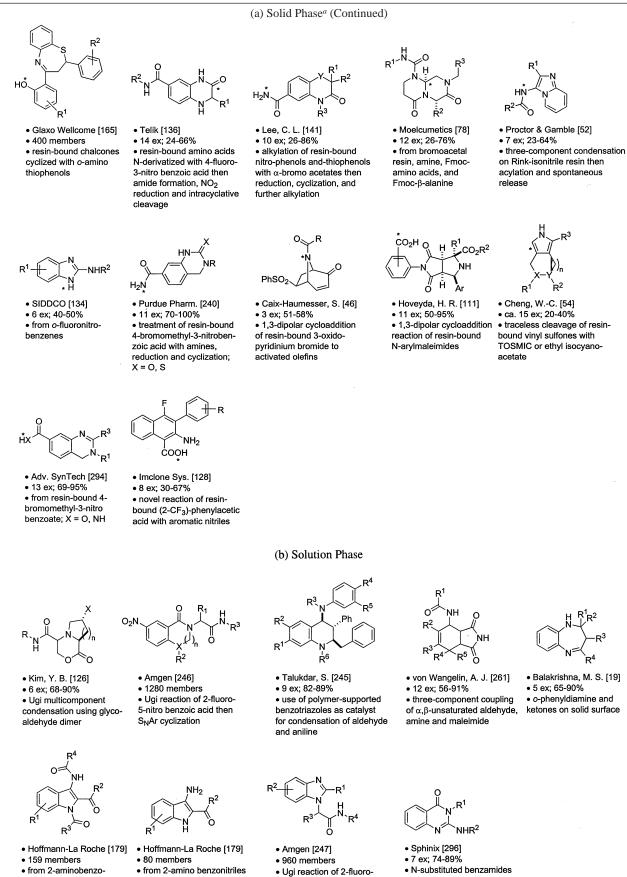
benzenes, mesylation, NO₂

reacted with o-fluoronitro benzenes, NO2 reduction, cyclization with BrCN then cleavage

• S_NAr of resin-bound 2-fluoroamines, NO2 reduction, cyclization with TCDI then cleavage

reaction of resin-bound triamines with thiocarbonyldimidazole and Hg(OAc)₂

Table 9 (Continued)



3-nitrobenzoic acid then

S_NAr cyclization

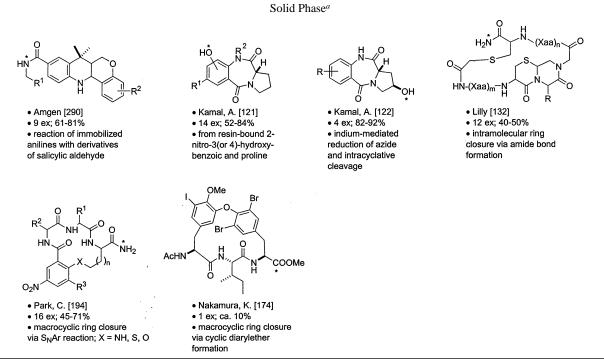
subjected to Kirsanov reaction

using polystyryl triphenyl-

phosphine resin

nitriles

Table 10. Polycyclic and Macrocyclic Synthesis



^a The asterisk (*) represents point of attachment to the resin.

Table 11.	Resin-Bound	Reagents	and Scavengers	Reported in 2001

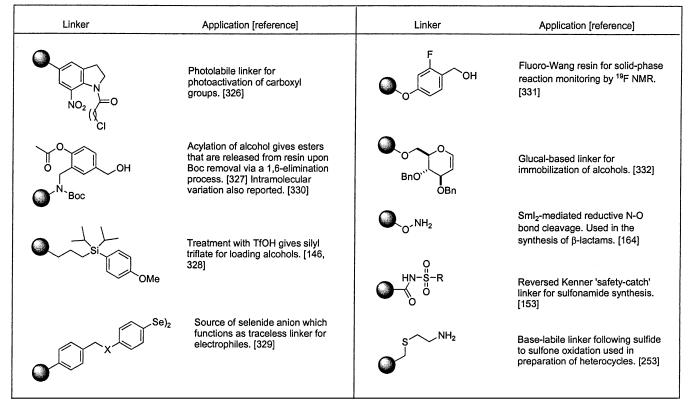
Reagent	Application [reference]	Reagent	Application [reference]
SeSO ₂ Ar	Vinyl sulfones from olefins. [208]	PPh ₂	Resin-bound iminophos- phoranes from anilines; preparation of quinazolines. [296]
O O CI SeBr	Chiral ethers from olefins. [254] Esterification of carboxylic acids. [286]	P(cHex) ₂	Pd-catalyzed aminations and Suzuki reactions. [196]
€ C C C C C C C C	Formatino of resin-bound aryl triflates from phenols. [192, 193]		Co-oxidant in the TPAP oxidation of alcohols. [333]
SO ₃ Sc(OTf) ₂	Nucleophilic addition to <i>N</i> -acylimino esters. [131]		Oxidation of alcohols to alehydes and ketones. [171, 229]
	RCM with electron deficient olefins. [210]	- chlorite	Oxidation of aldehydes to acids. [244]

Table 11 (Continued)

Reagent	Application [reference]	Reagent	Application [reference]
	Vinyl sulfone as Diels- Alder diene precursor. [54] Rink-isonitrile resin for multi-component		(a) Ureas from phosgene and amines. [191]; (b) Tetrahydroquinolines from aldehydes and aromatic amines. [245]; (c) Comparison to related triazole and benzotriazole leaving groups. [124]
€	reactions. [52] Purification-free method for the peparation of homoallylic alcohols. [13] Trityl isothiocyanate resin as precursor for resin- bound trityl thiosemicarbazide for preparation of heterocycles. [202]		O HN → N-Tf HN → NHBoc Traceless guanidinylating reagent for secondary amines to prepare N,N- disubstituted guanidines. [287]

Table 12. Solid-Phase Linkers Reported in 2001

Linker	Application [reference]	Linker	Application [reference]
NH	Phenanthridine-based linker. Release of carboxylic acids upon treatment with cerium ammonium nitrate (CAN). Stable to acids, bases, reductive amination. [318]	Si y	General synthesis of functionalized arylsilanes. [324]
	Preparation of dehydroamino acid amides. [319]		Linker that uses CAN as a cleavage reagent. Applied to the synthesis of β -lactams. [92]
FmocHN COOR	Resistant to basic and moderately acidic media, oxidation, and elevated thermal conditions. [320] Synthesis of Tryprostatin B. [262]	О С С С С С С С С С С С С С С С С С С С	Couples to acids to form ester linkage which can be cleaved by using both aqueous acidic and basic conditions. Intramolecular release to form hydantoins. [113]
O O O O S NH ₂	Synthesis of chiral amines in near quantitative yields in high enantiomeric purities. [321]		Synthesis of α-branched primary amines. [127] BOMBA resin for the traceless cleavage of heterocycles. [138]
Q X R	Synthesis of 1,2-diols via Pummerer cleavage strategy. [322]		
OH OtBu	'Safety-catch' ester linker. Linked esters are stable to nucleophilic chemistry. TFA-treatment gives the activated 2-hydroxyphenyl ester which readily reacts with amines to give amide cleavage products. [323]	(OC) ₂ Cr ² PPh ₂	Chromium carbonyl linker for attaching arenes to resin via π -bond ligand chemistry. [214, 325]



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