

Review

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J. Comb. Chem., 2005, 7 (6), 739-798• DOI: 10.1021/cc050082t • Publication Date (Web): 17 September 2005

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Volume 7, Number 6

November/December 2005

Reviews

Comprehensive Survey of Combinatorial Library Synthesis: 2004

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Received June 29, 2005

This is the eighth annual review in an ongoing series of comprehensive reviews in combinatorial chemistry highlighting developments in new methodology and synthesis of small molecule libraries.¹ The survey tallies 388 chemical libraries published in 2004,^{2–465} categorized according to biologically active libraries (Tables 1–5), libraries without disclosed biological activity (Tables 6–10), solid- and solution-phase reagents and scavengers (Table 11), linkers (Table 12), and polymer-supported chiral ligands (Table 13).

Affymax²⁵⁸ and Pharmacopeia^{153,154} independently reported the discovery of follicle-stimulating hormone (FSH) receptor agonists. The binding of FSH with its receptor constitutes a protein–protein interaction.

In addition to FSH agonist discovery, Pharmacopeia and their collaborators published the results of screening and optimization programs on glycine-2 transporter inhibitors⁴⁵⁷ and selective p38 kinase inhibitors.⁴⁶¹ The company is one of the original combinatorial chemistry companies founded in 1993, having licensed electrophoric-based encoding technology from Columbia University. Over the past decade, Pharmacopeia has amassed 7.5 million discrete small molecule compounds. The compound collection has been screened against dozens of targets, spanning all target classes: proteases, kinases, nonproteolytic enzymes, GPCRs, integrins, transporters, and other nonGPCRs. Diller and Hobbs⁴⁶⁷ published a statistical analysis of the company's historical high-throughput screening data examining the relationship between physical properties and substructures on the likelihood of compounds' displaying biological activity, irrespective of target class (structural properties

versus hit rate). Data mining revealed that rotatable bond count had the most significant impact on hit rate, with more rigid structures preferred. The optimal rotatable bond count was 6. Structures possessing functional groups capable of making strong intermolecular interactions were also favored. Compounds devoid of an amide bond were 2-fold more likely to be biologically active than compounds with just a single amide bond. Averages for molecular weight, lipophilicity, and numbers of hydrogen-bond donors and acceptors were in the typical range for drug molecules.

Fragment-based approach was used to identify modulators of metabotropic glutamate receptor 5.²⁴³ In a similar vein, a "directed" fragment-based approach furnished growth hormone secretagogue receptor (GHS-R) antagonists.⁴⁴⁸

A computational approach to the design of small molecule ligands for GPCRs activated by positively charged peptide ligands was described by Neurocrine Biosciences.²²⁷ The Pharmacopeia group published a review on library design strategies for targeting kinases and GPCRs.⁴⁶⁸

NeoGenesis honed an affinity selection-mass spectrometry (AS-MS) method for ligand identification.⁹ A protein is exposed to a collection of compounds, and any protein—ligand complex formed is separated by rapid, low-temperature, microscale size-exclusion chromatography. Dissociation of the ligand in a subsequent step and mass spectral analysis identifies its structure. As proof-of-concept, the methodology was applied to the discovery of dihydrofolate reductase inhibitors. Merck, in collaboration with NeoGenesis, identified a novel BACE-1 inhibitor from a multimillion compound library using AS-MS.⁸⁹

Diversity-oriented synthesis (DOS) libraries prepared by split-pool and parallel synthesis were reported by the

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Schreiber group.^{55,212,221,248} These sophisticated libraries contain skeletally and stereochemically diverse small molecules earmarked as biological probes in chemical genetics. A quantitative method for measuring the relationship between compound diversity and hit rate (cellular activity) was described,^{212,469} as well as a strategy guide for preparing DOS libraries.⁴⁷⁰ Also from Harvard University, the first 65-member library of DNA-tethered macrocycles employing DNA-template organic synthesis (DTS) was published by D. Liu and co-workers.⁴⁵⁶ DTS methodology was developed for the synthesis of *N*-acyloxazolidines.⁴⁶⁶

A new safety-catch linker was developed permitting the release of aldehydes and alcohols via the Pummerer rearrangement.³⁵⁸ Perfluoroarylsulfonate linkers were shown to behave as triflate equivalents in Pd-catalyzed deoxygenation and cross-coupling reactions.^{62,323} Microwave irradiation of methanolic ammonia solutions of halides and sulfonates affords high yields of amine salts without significant dimerization.³⁴¹ Two new examples of an expanding repertoire of polymer-supported ionic liquids as catalysts for nucleophilic substitution²⁰⁹ and the Stetter reaction,²² and an ionic liquid-based AMEBA-type linker,¹⁰² were reported.

New developments in multicomponent reactions continue to appear in the literature. 6,10,24,32,98,106,109,146,148,174,215,231,256,309,310,424,433 One noteworthy development was the work of Pirrung and Sarma. 309,310 They discovered tremendous rate enhancements in the Passerini and Ugi reactions (with α - and β -amino acids) when carried out in water at room temperature. Small demonstration libraries were synthesized. Products were isolated by simple filtration of the precipitated compounds. A new reagent, MP-glyoxylate, was shown to be a formal-dehyde equivalent in the 3-CC of imidazoheterocycles. 256

Last, Taddei collected and reviewed experimental procedures for functional group color tests on solid support,⁴⁷¹ and new self-indicating amine scavenger resins⁸¹ and pHindicating resins¹³⁵ were described by Bradley.

Methionine Aminopeptidase-2 (MetAP2) Inhibitors. MetAP2 is a potential molecular target for angiogenic-based diseases. The therapeutic rationale for this target is derived from studies with fumagilin, a naturally occurring irreversible inhibitor of MetAP2 displaying antiangiogenic properties. TNP-470, a semisynthetic analogue of fumagilin, inhibits angiogenesis and tumor growth in vivo. MetAP2 is a metalloenzyme that removes the N-terminal initiator methionine residue of nascent proteins. Bestatin and amastatin, 3-amino-2-hydroxyamides, are known reversible inhibitors of MetAP2 and its isoenzyme, MetAP1. Compound 1 possessing the methionine side chain inhibits MetAP2 with a $K_i = 5.7 \,\mu\text{M}$ and MetAP-1 with a $K_i = 63 \,\mu\text{M}$ (Figure 1). X-ray crystallography analysis of MetAP2 reveals its S1' binding pocket can accommodate larger P1' side chains, which fortuitously also trend toward greater selectivity relative to MetAP1. This is exemplified by the P1' cyclohexylmethyl-containing inhibitor **2**: $K_i = 0.6 \,\mu\text{M}$, MetAP2; $K_i = 84 \ \mu M$, MetAP1 (140-fold selective). Sheppard and co-workers at Abbott constructed library 1.2 (3), keeping P1' constant and exploring alternative functionality for the P2' alanine residue in 2.361 This was carried out by solutionphase parallel synthesis in which the corresponding Boc-



Figure 1. Methionine aminopeptidase-2 (MetAP2) inhibitors.³⁶¹

protected 3-amino-2-hydroxy acid was coupled with over 500 amines (amino acids and non-amino acids) via its activated HOAT ester. The SAR obtained revealed a preference for hydrophobic functionality, for example, halogenated arylalkyl and naphthylalkyl. In particular, naphthylamide 4 showed a 4-fold increase in binding affinity for MetAP2, albeit with somewhat diminished selectivity (75-fold) versus dipeptide 2. Follow-up library 1.3 (5) was then prepared in which the P1' side chains (\mathbb{R}^1) and amines (\mathbb{R}^2) were combined in a 2-dimensional array. In this instance, inhibitory potency and selectivity were found dependent upon both R¹ and R². The SAR obtained from the follow-up library was not immediately interpretable. A series of enzyme-inhibitor complexes solved by X-ray crystallography indicated significant movement of the enzyme's nonactive site residues accommodating variation of those portions of the inhibitor structure protruding out of the active site. Combinatorial synthesis was credited with optimization against unique and highly variable binding modes, affording MetAP2 inhibitors 6 ($K_i = 190$ nM) and 7 ($K_i = 20$ nM), each displaying >500-fold selectivity versus MetAP1.

Optimization of Cyclin-Dependent Kinase 5/p25 (ckd5/ p25) Inhibitors. The formation and deposition of cytotoxic neurofibrillary tangles in the brain is a hallmark of AlzReviews



Figure 2. Stepwise optimization of 2-aminothiazole inhibitors of cdk5/p25.¹⁶⁴

heimer's disease etiology. One mechanism by which this may occur is through hyperphosphorylation of tau protein by the anomalous cdk5/p25 complex. Inhibitors of this enzyme are therefore of interest as potential therapeutic agents for the treatment of Alzheimer's disease and other neurodegenerative disorders. High-throughput screening for inhibitors of cdk5/ p25 complex at Pfizer turned up thiazole 8 (Figure 2).¹⁶⁴ Its submicromolar affinity (IC₅₀ = 321 nM), low molecular weight (212 daltons), and chemical tractability made 8 a promising candidate for optimization. Thiazole 8 showed a high degree of selectivity against a broad range of kinases with the exception of cdk2/cyclin E, for which 8 was equipotent (IC₅₀ = 318 nM). High-throughput synthesis was used to carry out optimization, which was performed in four cycles (collectively library 2.3). In optimization cycle 1, the 5-isopropyl-2-aminothiazole was conserved and the amino group derivatized, yielding collections of amides and ureas (9). This led to phenylacetamide analogue 10 with a \sim 4fold increase in affinity (IC₅₀ = 64 nM). In the next round of optimization, a set of commercially available and customprepared 5-substituted 2-aminothiazoles were acylated with phenylacetyl chloride to yield collection 11. Keeping the phenylacetamido group constant gave rise to the 5-cyclobutyl analogue 12 with further increased binding affinity (IC₅₀ = 25 nM) and selectivity (3-fold). Larger cycloalkyl, alkyl, and phenyl groups at the 5-position decreased potency. Dimethylamino, which is sterically similar to isopropyl, but more polar, also resulted in loss of potency. Following the discovery of the 5-cyclobutyl as the optimal 5-substituent, two further optimization cycles took place, expanding the number of amide (optimization cycle 3; $12 \rightarrow 13$) and urea derivatives (optimization cycle 4; $13 \rightarrow 14$). The SAR revealed a preference for fused heteroaryl rings, for example, 15-17. Inhibitor 17 demonstrated a 60-fold improvement in potency at cdk5 (IC₅₀ = 5 nM) and a 10-fold selectivity over ckd2. Because only 2 of the 29 conserved ATP binding pocket residues in cdk5 and ckd2 are different, identifying highly selective ckd5 inhibitor is a demanding optimization exercise.

Inhibition of ATPase Activity of Human Papillomavirus 6 (HPV6) E1 Helicase. Infections caused by human papillomavirus result in a range of unpleasant conditions, including plantar and genital warts; laryngeal papillomatosis, a respiratory tract infection; and life-threatening cervical cancer. There are currently no marketed anti-HPV agents. The circular DNA genome encodes for 10 proteins (E1–E8 and L1, L2), which are all common to the >80 types of HPV. Of particular interest is the enzyme E1 (DNA) helicase, a highly conserved HPV protein. Inhibition of the E1 helicase should result in the inhibition of HPV replication, and therefore, the enzyme is considered an attractive molecular target for antiviral intervention. A high-throughput screen, measuring ATPase activity of recombinant HPV6 E1 helicase, was developed at Boehringer Ingelheim.¹²⁴ Evaluation of the company's compound collection resulted in the discovery of ((4-phenyl)phenylsulfonyl)acetic acid 18 as a reversible micromolar inhibitor of the enzyme (IC₅₀ = 2μ M; Figure 3). Acid 18 was subjected to a two-part hit-to-lead campaign. Part one explored the SAR of the sulfonylacetic acid and phenyl ring regions of the screening hit. This was carried out by traditional synthesis. An extensive survey of carboxylic acid isosteres and modifications to the sulfonyl group (>12 analogues) resulted in significant loss of activity, for example, 19 and 20. Similar exploration of the aryl ring revealed that the biaryl ring was required for activity (18 \rightarrow 21, 22), wherein the 4-phenyl ring tolerated substitution at the *meta* and *para* positions $(18 \rightarrow 23, 24)$. With this nascent SAR in hand, part two of the hit-to-lead exercise was initiated. Parallel synthesis was employed to further define the scope of 4-phenyl substituents with the goal of increasing the affinity of 18. Synthons 25a,b and 26a,b were synthesized and coupled with a diverse set of commercially available amines to generate amide library 2.29 (27: ~ 100 analogues total). Increased potency was achieved within the libraries with IC₅₀ values ranging from 400 nM up to >80 000 nM. The most potent analogues were the metasubstituted series of primary amides bearing a hydrophobic aromatic residue. Representative of the SAR were 29 (IC_{50}

Part 1 - Hit-to-lead defining SAR in regions (A) and (B):



Figure 3. Hit-to-lead for HPV6 E1 helicase inhibitors.¹²⁴

= 420 nM) versus **31** (IC₅₀ = 1300 nM) and **30** (IC₅₀ = 87 nM) versus **32** (IC₅₀ = 510 nM). Continued focused library synthesis of ~150 additional *meta*-substituted analogues (library 2.30 (**28**)) furnished compounds **33** (IC₅₀ = 9.2 nM) and **34** (IC₅₀ = 4.3 nM), with the latter analogue demonstrating a 500-fold improvement in potency versus the screening lead **18**. One issue with the series was the lack of cell potency and the propensity for the sulfonylacetic acid to decarboxy-late to the inactive methyl sulfone **35**. For this reason, sulfonacetic acid surrogates were revisited using **34** as a starting point. Unfortunately, none could be identified which maintained nanomolar potency. Two moderately active surrogates synthesized were **36a** (IC₅₀ = 630 nM) and **36b** (IC₅₀ = 740 nM). Neither of these compounds demonstrated whole-cell-based activity.

Affinity Selection-Mass Spectrometry: Identification of DHFR and β -Secretase Inhibitors. NeoGenesis reported an optimized automated process for identifying small molecule ligands from mass-encoded libraries on the basis of affinity selection-mass spectrometry (AS-MS; Figure 4).⁹ In this process, soluble protein (enzyme, receptor) is incubated with a compound library. The mixture is subjected to automated microscale size-exclusion chromatography (SEC) conducted at 4 °C in less than 30 s, separating any newly formed protein-ligand complex(es). The combination of speed and low temperature maintains the integrity of the protein-ligand complex, even for weakly bound ligands (K_d < 10 μ M) and moderate off rates ($k_{off} < 0.1 \text{ s}^{-1}$). The separated complex is directly routed to a reversed-phase chromatography column (60 °C, pH <2) where proteinligand dissociation occurs. The unbound ligand is then eluted into an electrospray HRMS for analysis. Because the libraries are constructed to minimize isobaric compounds by judicious choice of building blocks, data-searching algorithms can be used to identify the ligand by its molecular weight. Final confirmation of ligand structure is secured by independent synthesis of the ligand. If positional isomers exist (mass redundancy), then a sublibrary of ligands is synthesized. Highlights of the process include (1) no foreknowledge of the structure or biochemistry of the protein target is required; (2) immobilization or modification of the protein is unnecessary; (3) only submilligram amounts of purified, soluble protein is consumed (<1 μ g protein per 2500 compounds); and (4) screening can be conducted at a rate of $> 250\ 000$



 $.R^{1}$

Figure 4. NeoGenesis's affinity selection-mass spectrometry method for ligand identification.⁹

compounds/day employing mixture libraries prepared by solid- or solution-phase chemistry.

By way of showcasing the AS-MS process, a 2560member library was screened against *Escherichia coli* dihydrofolate reductase (DHFR). The library consisted of five 500-member libraries, including library 2.23 (**39**), prepared from bifunctional template **37** via acylation then epoxide ring opening ($37 \rightarrow 38 \rightarrow 39$) with 15 amines (same inputs, twice). The AS-MS process was conducted by incubating the 2560-member library at a 2.5 mM cumulative compound concentration with 5 μ M DHFR (2 μ L final volume, pH 7.5, phosphate buffer containing 2.5% DMSO, 100 mM NaCl) for 30 min at 37 °C, then chilled to 4 °C for SEC separation and further processing. In this way, 2 pmol of each library member (at $1.0 \,\mu$ M/component) and 10 pmol $(0.18 \ \mu g)$ of protein were used in the analysis. Screening vielded a single monochlorinated DHFR ligand at m/z 515.24. The ligand corresponded to (*R*,*S*)-40 or its positional isomer. These were the only two ligands in the 2560-member library that came within 0.05 amu of the measured molecular weight. Compound (R,S)-40 and its positional isomer were individually synthesized employing the same experimental method used to produce the library. By comparison of the molecular fragmentation patterns of the authentic samples versus the isolated ligand, the structure of the DHFR ligand was determined to be the structure, as indicated. The R- and S-isomers were then separated by chiral HPLC. (S)-40 was found to be the more active compound ($K_d = 15 \,\mu\text{M}$) versus its antipode (R)-40 ($K_d = 40 \ \mu M$). (S)-40 inhibited the growth of *E. coli* at an IC₅₀ of 29 μ g/mL.

In collaboration with NeoGenesis, Merck applied the AS-MS technology in a screening campaign to identify novel leads against β -secretase (BACE-1).⁸⁹ This resulted in the identification of structure 41 as the only compound emerging from a multimillion-compound library 1.8 (generic library structure was not defined). The nonpeptide BACE-1 inhibitor $(IC_{50} = 25 \ \mu M)$ was subjected to a round of optimization. Replacement of the N,N-diallylcarboxamide with a benzyl sulfonate $(41 \rightarrow 42)$ resulted in a ~17-fold increase in enzyme affinity. The stereochemistry present in 42 was critical for activity: (S)-42, $IC_{50} = 1.4 \ \mu M$ versus (R)-42, $IC_{50} > 100 \ \mu$ M. An X-ray crystal structure of the corresponding enzyme-inhibitor complex revealed that (S)-42 occupies the enzyme's S_4-S_1 subsites. Interestingly, the inhibitor makes no direct contact with the catalytic aspartic acid residues, but rather, its oxyacetamide NH forms a hydrogen bond with a catalytic water molecule situated between the aspartyl triad.

Positive Allosteric Modulators for mGluR5. The binding of glutamate and subsequent activation of the metabotropic glutamate receptor 5 (mGluR5) is coupled to increased function of the NMDA receptor. Suboptimal functioning of the NMDA receptor is associated with psychosis, including schizophrenia. Compounds that potentiate the binding of glutamate to mGluR5, that is, positive allosteric modulators, may therefore restore hypofunctioning NMDA receptors, leading to a new class of antipsychotic agents. A highthroughput FLIPR assay was developed at Merck to identify compounds with this biological profile.²⁴³ One compound to emerge from the screen was phthalimide 43: $EC_{50} = 740$ nM (Figure 5). This compound demonstrated a 6-fold potentiation of the response of CHO cells transfected with human mGluR5 to a low concentration of glutamate, the native agonist. The compound did not display agonist activity in the absence of glutamate, further confirming it as a positive allosteric modulator of the receptor. Lindsley led the effort to develop a SAR. In this regard, an iterative library approach to analogue synthesis was carried out, and some 1000 analogues (library 3.19) were reportedly prepared. Despite

Initial lead:



Figure 5. Positive allosteric modulators for mGluR5.²⁴³

the large number of analogues, the screening hit proved intractable. Compound 44, in which chlorine conservatively replaced bromine in 43, was one of only two structures (the other compound in which 2-pyridyl was replaced with 2,2hydroxyphenyl) that retained a reasonable level of potentiation. A new lead structure was required. To this end, a fragment-base library approach in which the picolinoyl amide was retained was adopted. Picolinoyl chloride 45 was then reacted with a diverse set of commercially available and inhouse heterocyclic amines 46 to give library 3.20 (47). The size of the library was not disclosed. This exercise led to the discovery of a new lead, 1,3-diphenylpyrazole 48: EC_{50} = 290 nM, \sim 3-fold potentiation. Three focused libraries (50) were then prepared. Commercially available 5-amino-1,3diphenylpyrazole 49 was derivatized with a range of carboxylic acids and chlorides. It was found that the phenyl derivative 51 was 7-fold more potent than 48. Activity was dependent on the substituent pattern in the phenyl ring $(EC_{50}$'s for ortho- (52), meta- (53), and para-toluyl (54) were $>10\ 000,\ 90,\ and\ 30\ nM,\ respectively)$. Analogue 55 (EC₅₀ = 10 nM, 4-fold potentiation) was selected for further



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Figure 6. Abbott's growth hormone secretagogue receptor antagonists.448

evaluation. It was devoid of ancillary activity against a panel of kinases and receptors. The mGluR5 potentiator readily diffused into the CNS as measured by selected brain penetration experiments. In a critical in vivo proof-of-concept study, 55 dose-dependently reversed the rodent acoustic startle response following amphetamine administration. This is a well-characterized animal behavior model of sensorimotor gating in which antipsychotic drugs are similarly effective.

Growth Hormone Secretagogue Receptor (GHS-R) Antagonists. Administration of a peptide antagonist of GHS-R results in reduction of food intake and body weight gain in diet-induced obese mice, suggesting that GHS-R antagonists may be useful therapeutics for treating obesity. Isoxazole 56 was identified from a high-throughput screening campaign for GHS-R ligands at Abbott Laboratories (Figure 6).448 This attractive hit was a rather potent antagonist possessing an $IC_{50} = 130$ nM against the receptor. Although extensive SAR led to more potent analogues, as a class, these displayed poor pharmacokinetics and were largely devoid of oral activity, the desired route of administration. Part of the problem was thought to be facile hydrolysis of the amide group in vivo. The isoxazole ring was replaced with a variety



Figure 7. Pharmacopeia's FSH receptor agonists.¹⁵³

of aryl and heteroaryl rings ($56 \rightarrow 57$). On balance, these analogues retained receptor affinity. This led to the hypothesis that the isoxazole ring served as a spacer element providing a preferred distance and dihedral angle between the aryl rings in 56. Ten scaffolds 58 were synthesized to test the hypothesis, and a new tetralin template, exemplified by 59, was found. The IC_{50} of 59 was 200-fold less potent $(IC_{50} = 2700 \text{ nM})$ versus the original hit (130 nM). Amide hydrolysis (metabolic instability) was still of concern in the new structure, and thus, the carbon atom α to the amide was quaternized by introducing alkyl, carboxylic, and assorted amine derivatives. This afforded the Boc-protected α -amino analogue 60, restoring receptor affinity of the original hit. Library 3.15 (61) was then prepared to uncover isobutyl as the optimal carbamate: 62, $IC_{50} = 11$ nM. Further SAR studies led to 7-methoxy analogue 63, $IC_{50} = 16$ nM. Importantly, compound 63 was orally bioavailable in the rat (F = 19%), with reasonable clearance, and displayed excellent specificity (no off-target activity) and low hERG channel binding. Tetralin 63 was a functional antagonist of GHS-R. Receptor affinity of the enantiomers was not reported, nor was 63's ability to elicit reduction of food intake in an in vivo model of obesity.

Encoded Libraries and the Discovery of Follicle-Stimulating Hormone (FSH) Receptor Agonists. FSH is 38-kD heterodimeric protein which binds to FSH receptor, a G-protein-coupled receptor (GPCR), setting in motion a signaling cascade leading to the growth of ovarian follicles and improved fertility. Purified or recombinant FSH is a biopharmacetical used to treat low fertility in women. FSH must be administered by subcutaneous or intramuscular injection, and an orally active agent is highly desirable. Because the binding of FSH with its receptor is in essence a protein-protein interaction, the discovery of small molecules is a challenge. This past year, research groups at Pharmacopeia^{153,154} and Affymax²⁵⁸ independently reported potent nonpeptide FSH receptor agonists. In both instances, screening hits came from large chemically encoded libraries.

Guo and co-workers at Pharmacopeia collaborated with scientists at Organon to discover FSH agonists (Figure 7).¹⁵³ Two million compounds constituting 37 structurally distinct ECLiPS libraries (split—pool libraries encoded with electrophoric tags) were evaluated in a high-throughput screen using a CHO cell line expressing human FSH receptor and the cAmp-response-element (CRE)-luciferase-reporter con-

struct. One of the 37 ECLiPS libraries may perhaps have contained the privileged biaryl motif and, hence, yielded (S)-64 and 65 as micromolar hits. A follow-up encoded library was synthesized to expand the nascent SAR. Thirty-one reaction vessels containing TentaGel resin 66 modified with the photolabile linker and encoded with a set of electrophoric halogenated tags were reacted with one of 31 primary amine inputs. Resin 67 so obtained was pooled and divided into 11 reaction vessels and acylated with one of 11 iodoaryl carboxylic acids and then tagged to give acylated resin, 68. It was subsequently pooled and mixed, and approximately one-third of the resin was divided into 17 portions and subjected to Suzuki coupling with one of 17 arylboronic acids, yielding 17 sublibraries of 341 compounds (70a). The remaining two-thirds portion of resin was divided thrice; coupled to *o*-, *m*-, and *p*-formylphenylboronic acids; tagged; pooled; mixed; and split into 25 reaction vessels, $68 \rightarrow 69$. Final resin treatment included a combination of reductive amination and N-capping, generating 25 sublibraries of 1023 members each (70b). The total number of library 3.11 compounds = 31 372. Compounds were cleaved from resin by photolysis after bead arraying. The structure identity of any given active was determined by ECGC analysis of the haloaromatic alcohol tags detached from beads in a separate tag-reading step and then confirmed by compound resynthesis. It was reported that 72 distinct structures with an EC_{50} < 10 μ M were found. One of the reductive amination sublibraries contained 25 actives and was analyzed with respect to distribution frequency of synthons. Interestingly, this analysis revealed a remarkable combinatorial dependence on synthon pairs $R^3 - R^2$ and $R^3 - R^1$. The frequency of R^3 ortho-compounds was tightly paired with acylic carbamates, as found in 65. The frequency of R^3 -meta-compounds were coupled with the cyclic scaffolds, represented by azepinone 71 and diketopiperazine 72. Last, R³-para-compounds were nearly exclusively associated with the hydantoin scaffold, represented by 73.

The diketopiperazine biaryls **64** and **72** obtained from the ECLiPS collection and follow-up library 3.11 were subjected to a second optimization step (Figure 8).¹⁵⁴ Parallel synthesis was used to prepare an additional ~300 compounds (library 3.12). Chemistry was developed to permit intracyclative cleavage of resin-bound Boc-protected amino esters to give the desamide biaryl diketopiperiazines that were not contained in the original ECLiPS library due to the synthetic strategy employed. The desamides **74** ((2*S*, 5*S*); EC₅₀ = 13 nM) and **75** ((2*S*, 5*R*,*S*); EC₅₀ = 1.2 nM) gave leads with greatly enhanced (>150-fold) potency over the original screening hit **64**.

In the Affymax study led by Maclean and Holmes,²⁵⁸ broad screening of the Affymax libraries afforded thiazolidinone **76** (EC₅₀ = \sim 5 μ M) as a hit structure (Figure 9). An encoded split—pool library 3.10 (**77**), employing alkylamine-based tags, was prepared. The library was composed of 35 R¹ amino acids × 35 R² aldehydes × mercaptosuccinic acid × 35 HNR³R⁴ amines. The last step was not pooled, but kept as separate sublibraries. The total number library compounds was nominally 42 875, but is significantly much larger, considering that it is a mixture of diastereomers. In-process





Figure 8. Optimization of Pharmacopeia's FSH receptor agonists.¹⁵⁴

and final library quality control indicated a synthesis success rate of 96%. The library was designed to be in part a followup on **76** and in part random for screening against future molecular targets. As a result, only 10 building blocks per input set were either identical or closely matched functionality in **76**. The remaining 25 building blocks per input set were randomly selected, in part, on the basis of their expected chemical performance. Hence, only $\sim 2\%$ of the library contained direct analogues of **76**.

Library 3.10 (77) was initially surveyed to identify active sublibraries. This was done by arraying the entire library, 35 sublibraries, in 96-well plates at 30 beads per well in DMSO, followed by full photolysis (1 h, 365 nm) and screening (FSH receptor reporter assay). Two active sublibraries were found. Two screening strategies were then employed to identify biologically active single beads. In the first strategy, so-called tiered release, 0.75 library equivalents (defined as the number of library beads equal to the number of compounds in the library) was arrayed at 10 beads per well (96-well plate) and briefly photolyzed (2 min, 365 nm)



Figure 9. Affymax FSH receptor agonists: library synthesis.²⁵⁸

to release half of the compound from the bead. Only one of the two sublibraries confirmed activity. The beads from the active sublibrary were rearrayed as single beads per well and subjected to a second photolysis treatment, releasing the rest (remaining half) of the compound. Following screening and decoding, the structures of submicromolar FSH agonists were revealed as **81** and **82**.

In the second screening strategy, 3 library equivalents of just the confirmed active sublibrary was arrayed in six 864well plates (high-density format) as single beads. Full photolysis and screening gave 63 active wells out of 5000 wells. Structures of the active wells were decoded, affording compounds 83 (EC₅₀ = 32 nM) and 84 (EC₅₀ < 100 nM), the former compound being significantly more potent than the original hit 76. Compound 83 differs from 76 only by a benzyl versus methyl substituent in the aryl ring. Interestingly, the original hit 76 was not found by either screening strategy. It was subsequently shown that 76 was sensitive to TFA used in the final library treatment (R¹ amino acid side chain deprotection). Surprisingly, 83 was apparently insensitive to TFA. Omitting this TFA-cleavage step, 76 was found upon screening 2 library equivalents of the appropriate sublibrary in high-density format. This report aptly highlights the advantages and pitfalls of screening large, chemically encoded libraries.

Small Molecule Library for GPCRs Activated by Positively Charged Peptides. GPCRs for melanocortins, gonadotropin releasing hormone, bradykinin, melanin concentrating hormone, vasoactive intestinal peptide, galanin, orexin, and selected chemokines continue to receive much attention in the pharmaceutical industry. This is due to their involvement in a range of pathophysiological conditions, including reproductive disorders, pain, and obesity. These receptors constitute a subclass of GPCRs that are all activated by peptide ligands carrying an obligatory positively charged residue. Saunders and co-workers at Neurocrine Biosciences designed a non-peptide library test set (library 3.18 (93)) of 2024 members targeting this subclass of GPCRs (Figure 10).²²⁷ They first defined the 5-dimensional space occupied by 81 560 biologically active compounds, including marketed drugs, by calculating their BCUT metrics (DiverseSolutions software) for H-bond donor and acceptor, two metrics of polarizability and charge. Relying on cell-based methodology, each of the 5 axes was divided into 10 bins, partitioning the entire drug space into 100 000 individual cells. The actual number of cells occupied by the 81 560 ligands was 8506 cells, or $\sim 8\%$ of the total. At this juncture, Saunders introduced the concept of analyzing neighboring cells. Experience had taught that although potent compounds reside in a few cells or a single cell, progressively less active



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Figure 10. GPCR screening library.²²⁷



Figure 11. Safety-catch linker cleaved by Pummerer rearrangement.³⁵⁸

compounds are layered in neighboring cells. In 5-dimensional space, each occupied cell has $3^5 - 1 = 242$ neighbors, and for the 81 560 ligands, raising the total number of cells populated to 91 080 cells, or 91%. A similar analysis was carried out on 111 known nonpeptide ligands for positively activated GPCRs. Mapping the results showed that only 6% of the drug space (6437 cells) was occupied, including nearest neighbor cell count, which projected into three of five dimensions. With this analysis in hand, 19 scaffolds containing at least one basic nitrogen atom were subjected to virtual library derivatization and enumeration. A virtual collection of 90 000 000 compounds resulted, each possessing a basic nitrogen atom within the scaffold as the key pharmacophoric element. The five BCUT metrics were calculated for each virtual compound to assess its location relative to the space occupied by the 111 known nonpeptide ligands. Seven of the scaffolds, 85-92, were selected for actual library synthesis. The scaffolds were derivatized employing acylation, sulfonylation, ureidoation, reductive amination, alkylation, and amidation reactions. This furnished library 3.18 (93) containing 2025 discrete compounds. Evaluation of 93 against three GPCRs, melanocortin-4 receptor (MC4-R), gonadotropin-releasing hormone receptor (GRH-R), and melanin-concentrating hormone receptor (MCH-R), resulted in a 4.5- to 61-fold increase in hit rate, as compared to a random library set of comparable size. The structures and K_i 's of several hits, for example, 94 and 95, were revealed for MCH-R. In addition, the computational analysis suggested that \sim 7000 compounds would be necessary to establish the boundaries of this GPCR subclass and provide a comprehensive targeted screening library.

New Linker Chemistry. Li and co-workers developed a new safety-catch linker 96 cleaved by Pummerer rearrangement, releasing aldehydes and alcohols (Figure 11).358 4-Hydroxythiophenol 97 was coupled to Merrifield resin in a high loading routine 3-step reaction sequence: S-tritylation, resin etherification, and S-trityl deprotection $(97 \rightarrow 96)$. The safety-catch linker 96 reacts with various electrophiles, such as halides and sulfonate esters (NaH is required as the base) and epoxides (Et₃N as the base), or alcohols under Mitsunobu conditions, yielding a sulfide resin 98. The sulfide linkage is stable to a wide range of reaction conditions. To cleave the linker, resin 98 is first oxidized to the sulfoxide 99. A survey of oxidants found t-BuOOH/10-camphorsulfonic acid in DCM was optimal (12 h, 25 °C), avoiding overoxidation to the corresponding sulfone. The oxidation reaction was conveniently monitored by magic angle spinning gel-phase NMR. Initial attempts to effect Pummerer rearrangement with sodium acetate/acetic anhydride proved too harsh, because the high temperature required was incompatible with the polystyrene bead. Switching to trifluoroacetic acid anhydride (TFAA) solved this problem, leading to the α -trifluoroacetoxythioacetals 100 in high yield at ambient temperature (THF, $0 \rightarrow 25$ °C, 45 min). Treatment of **100** with Et₃N in EtOH (2 h, 25 °C) released the aldehydes 101. By adding NaBH₄ to the cleavage cocktail, a one-pot procedure was developed to yield alcohols 102 directly.

Safety-catch linker synthesis:



Nitrobenzyl linker **103** was conceived as a new safetycatch linker for solid-phase synthesis possessing both acid and base stability (Figure 12).²⁹³ This is a Wang-type resin whose acid stability is dramatically improved by virtue of the nitro substituent. It is activated by reduction of the nitro group to the corresponding aniline, followed by acylation or sulfonylation. Products are released upon exposure to mild acid. Linker **103** was prepared upon Mitsunobu coupling of



Figure 14. Perfluoroarylsulfonate linker for cross-coupling.^{62,323}







Figure 15. Polymer-supported ionic catalysts.





Figure 17. Solid-phase synthesis of secondary and tertiary amines.²⁹⁷

4-hydroxy-3-nitrobenzaldehyde 104 to 2-hydroxyethylpolystyrene 105 followed by aldehyde reduction with NaBH₄ in MeOH-THF (1:9). Resin loading (determined by elemental analysis of 103) was reasonable at $\sim 1 \text{ mmol/g}$. To demonstrate the linkers's robustness to acid, naltrexone 106 was coupled to 103 (Mistunobu reaction conditions) to give resinbound phenol 107. Exposure of 107 to TFA-DCM (1:1) or t-BuOK in THF resulted in <5% cleavage of naltrexone.Nitro group reduction was readily carried out under standard reaction conditions using SnCl₂·H₂O. Sulfonylation of the anilino group rather than acylation (to avoid competing O-acylation) gave activated resin 108. The linker was cleaved (TFA-DCM (1:1)), resulting in 50% recovery of 106. In a further demonstration of the utility of the safety-catch linker, resin-bound naltrexone 107 was subjected to Fisher indole synthesis (4 \times arylhydrazines, TFA-DCM (5:95), 25 °C, 1



Figure 18. Microwave-assisted primary amine synthesis.³⁴¹

h) and N-alkylation (4 × ArCH₂Br, *t*-BuOK, THF, 25 °C, 14 h; **107** \rightarrow **109** \rightarrow **110**). Tin-mediated nitro reduction, N-sulfonylation, and acid cleavage produced the 16-member library **111**. The chemistry was performed by the directed sorting split—pool method with IRORI MicroKans. Final products were purified by preparative reversed-phase HPLC to remove *N*,*O*-dialkylated byproducts generated during the *t*-butoxide alkylation step.

Silyl linkers have broad appeal in solid-phase synthesis. Encoded diversity-oriented synthesis (DOS) libraries originating from the Schreiber group rely almost exclusively on immobilizing alcohol building blocks via the silyl ether. The linker supports aryls where the silicon-aryl bond is subject to ipso substitution to yield traceless cleavage products. Cleavage conditions are tunable, depending upon the type of alkyl substituents present on the silicon atom. Takahashi and co-workers reported an efficient synthesis of a range of polymer-supported silvl linkers employing di-Grignard chemistry (Figure 13).¹¹⁰ Pentane-1,5-di(magnesium bromide) 112 (12 equiv) was coupled to benzyl chloride-bearing resins (113), including Merrifield, Argogel-Cl, Wang-Cl, and Argogel Wang-Cl, in the presence of CuBr•Me₂S. The bulky nature of the resin permitted selective monoalkylation of 113. After filtration to remove excess 112, the reactive intermediate Grignard resin 114 was further treated with solutions of chlorodiisopropylsilane and chlorodiphenylsilane, affording 115 and 116, respectively. For the Wang-type resins, the



Figure 19. Formaldehyde equivalent in 3-component condensation.²⁵⁶

sequential reactions proceeded at room temperature. Resin loading was established by the Fmoc cleavage test via chlorination (SiH \rightarrow SiCl) and attachment of *N*-Fmoc-2-aminoethanol. Loading was adequate, averaging 50–75% of the initial loading. The silane resins were activated to the corresponding silyl chloride and silyl triflate resins using standard methodology for the attachment of alcohol substrates and formation of silyl ethers **117**. A novel one-step dehydrosilation method was also developed to attach alcohols directly to the diisopropylsilane resin **115** using B(C₆F₅)₃.

Two research groups, one led by Cammidge⁶² and the other by Ganesan,³²³ independently developed perfluoroarylsulfonate linker 118 (X = O or NH) as a polymer-supported "triflate" equivalent (Figure 14). Their research disclosures appeared as back-to-back communications in Chemical *Communications*. The readily available acid chloride **119** was coupled to resin 120 via an ester or amide linkage, affording sulfonyl chloride resin 121. Resin 121 smoothly reacted with a series of phenols in high yield to give resin-bound sulfonate esters 118. Both research groups demonstrated the opportunities for Pd-catalyzed traceless cleavage. For example, the reductive cleavage by transfer hydrogenation gives deoxygenated products 122. Site-specific isotopic labeling of aromatic compounds was achieved by substituting formic acid with deuterium-labeled formic acid in the reaction. Ganesan reported a $\sim 10\%$ higher yield of products 122 using amide versus ester linked tetrafluorosulfonate.323 Resin 118 also performed well in the Suzuki–Miyaura (118 \rightarrow 123), Mizoroki-Heck (118 \rightarrow 124), and Negishi (118 \rightarrow 125)



Figure 20. Multicomponent reactions in water.^{309,310}

cross-coupling reactions. Perfluoroarylsulfonate resins have excellent shelf stability.

Polymer-Supported Ionic Catalysts and Ionic Linkers. Examples of polymer-supported ionic catalysis and ionic liquid-based linkers representing a new and potentially burgeoning family of reagents for high-throughput synthesis were described in 2004. Ionic liquids are recognized as alternative reaction media for conducting a variety of chemical transformations. They are documented to accelerate chemical reactions; act as catalysts; and in some instances, provide cleaner reaction products with reduced byproduct formation. Nucleophilic substitution reactions with alkalimetal salts proceed with greater efficiency in ionic liquids containing imidazolium cations. Due to the difficulty in extracting polar products from such reaction media, Chi and co-workers investigated polymer-supported imidazolium salts **126a**-c as catalysts for this reaction class (Figure 15A).²⁰⁹ The polystyrene-based ionic-liquid system 126a was synthesized by simply heating the chloroalkyl ether resin 127 with N-methylimidazole 128 (neat, 90 °C, 72 h). Counterion exchange afforded resin-bound imidazolium salts 126b and 126c. A survey of the new resins in a representative fluorination reaction identified **126b** as the superior reagent, which could be used catalytically. Displacement of a primary mesylate with CsF in acetonitrile in the presence of 126b



Figure 21. Schreiber's diversity-oriented synthesis (DOS) libraries.

(0.5 equiv) gave 100% conversion in 2 h at 100 °C versus <5% product formation without **126b**. Ionic resin **126b** also catalyzed the S_N2 reaction using a range of nucleophiles (**129** \rightarrow **130**).

High-loading ROMPgel-supported thiazolium iodide **133** was developed by Barrett (Figure 15b).²² This ionic ROMPgel proved to be an effective catalyst in the Stetter reaction, the umpolung condensation of aldehydes **134** with, in this case, an α , β -unsaturated aryl ketone **135**, to give 3-oxoketones **136**. Triethylamine was the preferred base to generate the ionic catalytic species **137**. The catalyst was recovered up to four consecutive times without compromising its activity. Some 17 condensations were reported. The yields ranged from 68% to 99%, and product purity was excellent.

Ionic liquids have extremely low vapor pressure, are stable to heat, and form a separate phase in the presence of both aqueous and organic solvents. It was this latter physical property that led de Kort and co-workers to prepare the ionic AMEBA-type linker **140** (Figure 16).¹⁰² Microwave heating **141** with *N*-methylimidazole **128** furnished imidazolium salt

140 in quantitative yield. Linker 140 formed a homogeneous solution in the ionic liquid, [bmim][PF₆], and underwent sequential reductive amination and acylation/sulfonylation employing standard reaction conditions $(140 \rightarrow 142 \rightarrow 143)$. Taking advantage of ionic-phase assisted extraction, excess reagents were removed after each step by washing the reaction solution with Et₂O, then water; intermediates 142/143 remained in the ionic liquid. Linker cleavage was carried out using 60% aqueous HPF₆ (10-min exposure). Although cleavage occurred readily with TFA, HPF₆ was the preferred acid to avoid generating mixed counterions of the ionic liquid. Amide derivatives 144 were obtained upon quenching the cleavage reaction with saturated aq NaHCO₃ and extraction with organic solvent. Product yields were comparable to those obtained using AMEBA resin.

Amine Synthesis. As part of a broader research effort to develop an expedient solid-phase synthesis of polyamines, Franzyk investigated the synthesis of secondary and tertiary amines via direct displacement of resin-bound halides (Figure 17).¹⁹⁷ Initial attempts focused on the displacement of

polystyrene *N*-trityl-linked 1-iodopropylamine with primary amines. The approach was abandoned when issues with onresin halide formation and cross-linking during the S_N2 reaction surfaced, this despite the use of lower loading strategies. Ultimately, a protocol was worked out whereby primary and secondary amino alcohols, immobilized via a *N*-carbamate on Argopore Wang resin (145 \rightarrow 146–149), were efficiently converted to iodides 150 (I₂/Ph₃P) and displaced with a variety of amines, including unprotected amino alcohols (10 equiv of a 1.0 M solution of amine in DMF, 50 °C, 6 h). Following resin cleavage, diamines 151 were isolated in good yield (50–88%) with purities >90%.

An atom economical synthesis of primary amine salts 153 from halides, tosylates and mesylates 152 was achieved by Saulnier using simple microwave irradiation of their solutions in methanolic ammonia (Figure 18).³⁴¹ The optimal substrate concentration was established as 0.04 M (0.25 mmol of substrate in 6.25 mL of 7 M ammonia in methanol) and irradiation at 100-130 °C from 0.25 to 2.5 h. Some 20 different substrates were studied, represented by 153-164. The yields of the amine salts were in excess of 90%; secondary amine byproducts were typically <4%. Workup requires only solvent evaporation. Lower concentrations of ammonia led to an increase in secondary amine byproducts. In one specific example studied, substrate 165 gave 96% yield of the amine salt 166 using the microwave-assisted conditions (100 °C, 15 min) with <4% dimer **167** formation. Conducting the same reaction at room temperature resulted in complete conversion of substrate to \sim 75% yield of 166 and $\sim 25\%$ yield of 167. Volatile primary amines, for example, $156 \rightarrow 162$, are accessible because it is their nonvolatile salts that are isolated from the displacement reaction.

Multicomponent Condensations. The application of multicomponent condensation reactions in combinatorial library synthesis is well-known,472 dating back to the pioneering work of Armstrong.⁴⁷³ This remains a very active area of research. The 3-component condensation (3-CC) of 2-aminoazines, aldehydes, and isonitriles yields 2-substituted 3-amino-imidazoheterocyles. Such heterocyclic systems are found in biologically active compounds, including marketed drugs. Kercher and Lyon at Array Biopharma²⁵⁶ found that glyoxylic acid, in solution or immobilized on macroporous polystyrene resin, MP-glyoxylate 168, serves as a formaldehyde equivalent in this 3-CC, affording 2-unsubstituted-3-amino-imidazoheterocyles (Figure 19). In solution, the 3-CC/decarboxylation proceeds at room temperature. Using reagent 168, the reaction is conducted at 50 °C in dichloroethane (DCE)-MeOH. A Lewis-acid catalyst is not required. The scope of the 3-CC reaction was investigated using a variety of 2-aminoazines (173–177 as representative products). 2-Aminopyridines and 2-aminopyrazines were preferred inputs in terms of yield and purity of isolated reaction products.

The rate of multicomponent condensation reactions is profoundly influenced by solvent. In an elegant study of the Passerini reaction, Pirrung and Sarma³⁰⁹ showed that substituting water for organic solvent resulted in a rate acceleration of 18-fold ($178 + 179 + 180 \rightarrow 181$; Figure 20). Further rate enhancements were seen with LiCl and glucose as additives. The reaction in water proceeds faster at 4 °C versus 50 °C. The hydrophobic effect and cohesive energy density of water are thought to be factors contributing to rate acceleration. Similar rate enhancements were observed in the Ugi reaction $(178 + 179 + 180 + 182 \rightarrow 183)$. The condensation of the β -ketoacid **184**, amine **182**, and isonitrile 185 in water yielded a β -lactam 186 (72 h, 65%). This reaction does not proceed in organic solvent. A small demonstration library of Passerini products 187 ($4 \times 4 \times 4$ = 32 members) was prepared by stirring an acid (10%) excess) and equimolar amounts of aldehyde and isonitrile in water for 3-6 h at 25 °C. Solid products were isolated by simple filtration, while liquid products were isolated by extraction. The average yield and purity of the library was 87%. Libraries of Ugi products **188** and β -lactams 189^{310} were similarly prepared using water as the reaction solvent.

DOS Libraries. Library synthesis in Schreiber's laboratories at Harvard continues to emphasize design strategies that furnish collections of skeletally and stereochemically diverse small molecules (Figure 21). Complexity-generating reactions coupled with appending chemistries are represented by the encoded furan-derived library $(190 \rightarrow 191a-e)$,⁵⁵ based, in part, on the Achmatowicz reaction, the encoded spirooxindole library **195**,²⁴⁸ utilizing a 3-component reaction, and parallel libraries of nine-membered biaryl rings 201²²¹ and pyranoside-derived bicycles 210.112 In the library collection 191, substrates possessing pre-encoded skeletal information, termed σ -elements,⁴⁷⁰ were combined combinatorially under a common set of reaction conditions. All possible combinations of building block and stereochemical and skeletal diversity were efficiently assembled. This is analogous to the process of protein folding in which primary amino acid sequences (" σ -elements") are transformed into diverse 3-dimensional structures. Synthetic methodology was described for constructing libraries of alkaloid-like compounds $204a - e^{390}$ and macrocycles 208.³³⁴ Skeletal diversity in 204 was achieved by manipulating resin-bound reactive dihydroisoquinoline and dihydropyridine intermediates. Statistical and computational analyses were carried out to understand the relationship between structure and cell activity.^{212,469} No other research group worldwide is operating at this level of sophistication with regard to library design, synthetic intricacy, and assembly. Other reports of DOS methodology and libraries include 4-oxo- and 4-chloropyrido[2,3-d]pyrimidin-7(8H)-ones,⁴⁶⁴ 1,2-disubstituted and 1,2,5-trisubstituted pyrroles,¹⁵⁸ tetrasubstituted olefins,^{458,459} C1-nitrogen iminocyclitols,²⁴⁴ azaspirocycles,⁴²¹ polysubstituted spirotriones,460 sulfotransferase inhibitors,34 and tetrahydro-1,2-oxazines.433

Acknowledgment. The author is indeed indebted to the continued support and dedication of Karen Rivera who rendered the extensive chemical structure drawing of structures found in the tables. Appreciation is expressed to Paul Tuthill who assisted in organizing and editing portions of this year's manuscript.

Table 1. Chemical Libraries Targeting Proteases^a



Enzyme: Plasmepsin I and II (*Plasmodium falciparum*) Activity: $K_i = 0.5 \text{ nM}$, Plm I; $K_i = 2.2 \text{ nM}$, Plm II

Table 1. (Continued)









^a Asterisk (*), point of attachment to resin.

Table 2. Chemical Libraries Targeting Nonproteolytic Enzymes^a







Enzyme: Thymidine kinase-2 Activity: IC_{50} = 19 μM







Enzyme: Dihydroneopterin aldolase Activity: $IC_{50} = 68 \text{ nM}$

Table 2. (Continued)





Enzyme: MTA nucleosidase

Activity: K_i = 2.8 nM



^{*a*} Asterisk (*), point of attachment to resin.

Table 3. Chemical Libraries Targeting G-Protein-Coupled Receptors^a

Alphabetical listing NO₂ NR³R⁴ Library: 3.1 Name: Sulfamoylbenzamide NO2 н Ś, Size: 71 members Reference: Ritchie, T. J.; et al. [325] 2R1RN ő

Receptor: Bradykinin-1 (BK-1) Activity: K_i = 239 nM



Receptor: CCR-2b Activity: IC₅₀ = 0.71 µM





Activity: EC₅₀ = 290 nM; ~3 fold potentiation



Table 3. (Continued)



Target: NMDA Activity: $K_i = 2 \ \mu M$ (NR2B-selective NMDA antagonist)

Library: 3.27 Name: Benzimidazole Size: ca. 100 members Reference: McCauley, J. A.; et al. [271]



Receptor: Prostaglandin D_2 (human) Activity: $IC_{50} = 13 \text{ nM}$





Integrin: $\alpha_{v}\beta_{3}$ Activity: IC₅₀ = 2.6 nM, $\alpha_{v}\beta_{3}$; IC₅₀ = 3850 nM, $\alpha_{IIb}\beta_{3}$

Table 4. (Continued)





Activity: IC₅₀ = 0.2 mM (antagonist)

CI

Table 4. (Continued)



Library: 4.20 Name: Amide analog Size: 300 members Reference: Lu, Y.; *et al.* [254]

Library: 4.21 Name: 1,3-Oxazolidine Size: Not defined Reference: Hahn, H. - G.; et al. [155]

Library: 4.22 Name: Not defined Size: 33,040 members Reference: Jiang, S.; *et al.* [192] Note:"Universal" library of "drug-like" compounds from ChemBridge Corp.



Commercial library

R^{N.}

H

Ń

ŅН

x=0

.R²



Target: HCV-internal ribosome entry site (HCV-IRES) Activity: IC_{50} = 12 μM



Target: ELKI luciferase assay Activity: IC₅₀ = 9 μ M; GI₅₀ = 16 μ M, HCT-116 cell proliferation assay



Target: $\beta\text{-Hexosaminidase}$ release from mast cells Activity: 47% inhibition at 3 µM

CI



Target: HIV fusion Activity: EC₅₀ = 1 µM, HIV-1_{IIIB} replication

Library: 4.23 Name: Arylpyrimidine Size: 24 members Reference: Wang, G. T.; et al. [410]



NН

OH.

ÇF₃

13 N= NH N=N Target: LFA-1/ICAM-1 Activity: IC50 = 51 nM

NH

OH.

Library: 4.24 Name: Tryptophan derivatives Size: *ca*. 58 members Reference: Burdick, D. J.; *et al.* [54]

Library: 4.25 Name: Triazine Size: 2688 members Reference: Uttamchandani, M.; et al. [398]





0

N

Target: ICAM-1/LAF-1 Activity: IC₅₀ = 0.25 μM

Table 4. (Continued) CI N CI Library: 4.26 N Name: Benzimidazole Size: ca. 23 members ŇН NН CI CI Reference: He, Y.; et al. [162] R Target: E. coli 16S ribosomal RNA A-site Activity: IC_{50} = 60 μ M Library: 4.27 Name: 2-Quinoxalinol analog Size: *ca*. 75 members Reference: Zhang, L.; *et al.* [440] \mathbb{R}^2 H R¹ х 0 HO N NH₂ HO N NH₂ Target: TNF- α release (mouse macrophage) Activity: IC_{\rm 50} = 0.4 μM NR¹R² N 0 0 0 Library: 4.28 Name: 1,4-Substituted naphthalene Size: 32 members Ö O Reference: Ji, T.; et al. [190] N Ő N Target: Transcription factor PBXI/DNA binary complex Activity: IC $_{50}$ = 65 μM Library: 4.29 Name: Pyridazinone Size: 25 members Reference: Gouault, N.; *et al.* [147] R¹ \mathbb{R}^2 Ň .N 0 N' 0 Ň R³ Target: Isolated rat aorta Activity: Emax = 70.5% vasorelaxation Library: 4.30 Name: Benzoxazine R 0 O R Size: 19 members Reference: Caliendo, G.; *et al.* [61] N ò Ô S N N Target: Isolated rat aorta Activity: 97.6% of relaxatin at 10 4 M (EC $_{\rm 50}$ = 41 $\mu M,$ vasorelaxant activity) Library: 4.31 Name: Polyphenylurea Size: 89,856 members Reference: Schimmer, A. D.; *et al.* [343] Note: Positional scanning library. R^2 0 II С N H R¹ R⁴ Target: XIAP-mediated derepression of caspase-3 Activity: $IC_{50} = ca$. 10 μ M (Jurkat luekemia cells) Library: 4.32 Name: Hexahydroindoloquinolizinone Size: Not defined Reference: Jennings, L. D.; et al. [189]

Target: ZipA-FtsZ protein-protein interaction (antiinfective target) Activity: 46% inhibition at 1 mM

C



Library: 5.3 Name: Duocamycin analog Size: 45 members Reference: Ham, Y.-W.; et al. [156] Note: Selection assay that enables the direct identification of the most effective DNA alkylating agents from a mixture library.



R³



Target: H-Ras transfected MDCK-F3 cells Activity: $IC_{50} = 10 \ \mu M$ (phenotype reversion)

Target: L1210 cells Activity: IC₅₀ = 3 pM

S

Library: 5.4 Name: Acylpiperazine Size: 320 members Reference: Gerlach, M.; et al. [140]

Library: 5.5 Name: Crambescidin analog Size: ca. 10 members Reference: Aron, Z. D.; et al. [12]







Activity: IC₅₀ = 20 nM



Target: *in vivo* tumors (assorted tumor cell lines) Activity: Active in solid tumor mice model





Microbe: *P. falciparum* Activity: IC₅₀ = 3 nM







Reviews

Table 6. (Continued)





• Olsen [297] ca. 8 ex; good yield
resin-bound (trityl)amino alcohols converted to iodides then displacement with amines



R8

H₂N



 Olsen [298] · 3 ex; good yield from resin-bound ethylenediamine via Fukuyama-Mitsunobu alkylation sequence













Table 8. (Continued)

using TsCI/NaOH



facilitated by polymersupported reagents

Table 8. (Continued)







methyl cyanoacetate; X = CI, O





isothiocyanates or amines

dehydration

and phenylvinyl sulfide; oxidation and aromatization

amino)-prop-2-enoates

Table 9. (Continued)



^a Asterisk (*), point of attachment to resin.



Part A: Solid-phase



Saha [338]
48 members from resin-bound amino acids, o-fluoronitrobenzene and o-nitrobenzaldehydes



Krishnan [221]
 202 members

 multi-step sequence using amino alcohol, aldehyde and bromophenol inputs







Chem [79]
12 ex; 50-72% Bischler-Napieralski reaction for intracyclative cleavage

٠Ś

Tenuecyclamide A and related analogs • You [432] • ca. 4 ex; 38-71%

 sequential coupling of heterocyclic amino acids on resin, cleavage and macro cyclization



 Nielsen [288]
 ca. 20 ex; high purity
 intramolecular Pictet-Spengler
 reaction using N-Boc-1,3-oxazinanes as masked aldehyde equivalents



 Kamal [197]
 6 ex; 64-72%
 reaction of resin-bound isatoic anhydride with proline then TFA-mediated cleavage



• Hoesl [169] · 8 ex; good purity one pot aza-Wittig/heterocyclization/substitution from resin-bound benzimidazoles



R

```
• Wang [413]
• 10 ex; 69-90%
```

 3-component aza-Diels-Alder reaction of PEG-supported benzaldehyde, anilines and dihydrofuran/pyran



Cironi [85] 2 ex; good yield
 multi-step sequence
 from resin-bound bisarylacetylene



 Kamal [197]
 6 ex; 64-72%
 reaction of resin-bound isatoic anhydride with proline then TFA-mediated cleavage



 Schmidt [344]
 24 members study exploring factors controlling macro-cyclization



Wang [413]
10 ex; 53-78%

 3-component aza-Diels-Alder reaction of PEG-supported benzaldehyde, anilines and dihydrofuran/pyran then DDQ



Cironi [86]
3 ex; 53-86% Sonogashira cross-coupling of resin-bound phenolic iodides and 5-chloro-7-ethynylquinolin-8-yl acetate

then base treatment and cleavage



 Kamal [199]
 8 ex; 57-65%
 DIBAL reduction of resinbound thioesters then intramolecular aza-Wittig



- Arya [15]
 100 members
 - resin-bound hydroxyindoline acylated with N-nosylamino acids then tricycle formed by intramolecular Mitsunobu

Table 10. (Continued)



^a Asterisk (*), point of attachment to resin.







• water-soluble protecting group for amines; removed with dilute base (5% aqueous NaHCO₃)

 activation of enol ethers and glycals, introduction and cleavage of THP ethers Pd(II) complex for Suzuki-Miyaura cross-coupling

 reaction with Grignard compounds, amines and alcohols to give immobilized acyl, amide and ester ylides





Table 12. Polymer-Supported Linkers



MPEG-OCH₂SCH₃ Oikawa [294]

· formylacetal linker on

soluble support



CI N





 Ohno [293] · linker stable to strong acid or base; upon nitro reduction and acylation it is cleaved with mild acid

0

 Boas [39] · backbone amide linker cleaved with mild acid



 Mogemark [277] synthesis of n-pentenyl glycosides

• Timmer [393] liberates cyclopent-2-yl alcohols via RCM; application in oligosaccharide synthesis



• Boas [39] · backbone amide linker cleaved with mild acid



 Lazny [228] · new triazine linkers for the direct generation of polmersupported diazonium ions



 Cammidge [62] solid-phase equivalent to the triflate group



 Revell [323] · reaction with phenols yields arylsulfonates for traceless synthesis and Pd-catalyzed cross-couplings



 Lober [250] REM resin prepared using "click" chemistry







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CC050082T