

Review

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Reviews

Comprehensive Survey of Combinatorial Library Synthesis: 2002

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The sixth annual comprehensive survey on combinatorial chemistry¹ records a total of 388 chemical libraries published in 2002,^{2–414} a 25% increase from the previous year. Libraries are divided into two major categories: those with disclosed biological activity (122 entries; Tables 1-5) and those without accompanying biological data (266 entries; Tables 6-10). The screened libraries are segregated by molecular target class: proteolytic enzymes (Table 1), nonproteolytic enzymes (Table 2), GPCRs (Table 3), non-GPCRs (Table 4), and cytotoxics/anti-infectives (Table 5). Scaffold derivatization and acyclic, monocyclic, bicyclic/spirocyclic, and polycyclic ring syntheses compose Tables 6-10, respectively. Each of the latter headings is further subdivided into solid and solution phase. Table 11 covers polymer-supported reagents and scavengers; Table 12 covers new solid-phase linkers, while a new Table (13) is introduced on polymersupported chiral ligands. There are 67 entries in Tables 11-13, 90% of which come from academic laboratories, indicating that this group continues to be the principal driver for new solid-phase methods development. Overall, academics publishing solid- and solution-phase research outpaced industry by a 2:1 margin. The percentage of published libraries synthesized using solution-phase techniques retreated somewhat from last year to \sim 30%, while the average number of compounds in any given biologically active library

remained relatively steady (75%, <100 members; 90%, <1000 members).

Iterative library design/synthesis strategies blending solidand solution-phase protocols in lead optimization increased significantly. Some 20% of the libraries in Tables 1-5employed multiple streamlined synthesis techniques, suggesting researchers are electing to use all combinatorial tools at their disposal as appropriate to solve optimization problems. Examples include procollagen C-protease inhibitors (library 1.8),¹⁰⁴ caspase-3 inhibitors (library 1.23),⁸⁸ p56^{Lck} inhibitors (library 2.4),⁷⁴ cyclin-dependent kinase-2 inhibitors (library 2.6),²⁰³ neurokinin-1 antagonists (library 3.16),⁴¹ oxytocin antagonists (library 3.24),³⁸⁸ and Src SH₂ ligands (library 4.18).^{110,111} Tandem resin-solution chemistry was employed to construct nitrile-containing libraries yielding inhibitors of dipeptidyl peptidase IV (library 1.18)367 and cathepsin S (library 1.21).374 A clinical candidate emerged from library 1.18.

Therascope AG applied dynamic combinatorial chemistry to identify novel neuraminidase inhibitors.¹⁶⁴ Treating a diamine template with a mixture of aldehydes in the presence of enzyme and borohydride resulted in the amplification of a very small number, from potentially thousands, of transient hemiacetal/imine species that were reduced to give inhibitors with micromolar affinity. A striking illustration of a "privileged" GPCR scaffold was reported by Merck.³⁸⁰ A 128 000member mixture library of 2-aryl indoles was active against

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many families of GPCRs (16 screens), from which a number of highly potent and selective ligands were found following library deconvolution. Four companies, Roche,⁷³ Merck,²²⁸ Celltech,²²⁹ and Genentech,⁶⁴ described libraries targeting VLA-4 ($\alpha_4\beta_1$ integrin). The first three companies started with structurally similar dipeptide antagonist leads possessing poor pharmacokinetic properties. Libraries synthesized to circumvent this problem met with varied results.

Several academic groups published accounts of diversityoriented synthesis (DOS).^{100,173,213,217,218,258,266,276,336} The encoded libraries of Schreiber and co-workers^{213,217,218,258,336} for application in chemical genetics are characterized by natural product-type complexity and strive to introduce structural diversity through synthetic "branching points" within the constructs. In one example alone, 18 library compound crystal structures were generated as visual documentation of topological inimitability.

With regard to new methodology, several research groups employed Katritsky's benzotriazole chemistry to create heterocyclic libraries;^{193,195,248,249,319,366} a phase-switch protocol for the clean conversion of primary amines to secondary amines was reported by Pellitier;²⁹¹ and Hlasta described a traceless, multicomponent solid-phase synthesis of 2-substituted azoles via transient resin-bound azolium ylides.¹⁰⁸ Lindsley created an amide-containing series of fluorous scavengers that, surprisingly, were not retained on the standard Fluoro*Flash* SPE column alone, but required an ion exchange precolumn to fully remove the novel scavengers from reaction mixtures.²³³ These and other combinatorial chemical highlights are detailed below.

Nitrile Libraries as Inhibitors of Serine and Cysteine Proteases. Dipeptidyl peptidase IV (DPP-IV) is a serine protease with high substrate specificity cleaving N-terminal dipeptides Xaa-Pro or Xaa-Ala 1 from regulatory polypeptides (Figure 1). In vivo, DPP-IV inactivates glucagon-like peptide-1 (GLP-1), the most potent of the insulinotropic hormones. Selective inhibitors of DPP-IV have been shown to control blood glucose by extending the duration of action of GLP-1 to stimulate insulin secretion, inhibit glycagon release, and slow gastric emptying. In 1999, researchers at the Novartis Institute for Biomedical Research announced the slowly binding DPP-IV inhibitor 9 as a clinical development candidate for type-2 diabetes. Details of the discovery effort were not presented. This past year, it was revealed that combinatorial chemistry played a key role in its discovery.³⁶⁷ Previous inhibitor design strategies were based on L-amino acids with a protonatable N-terminal primary amine in the enzyme's P2 site. Novartis researchers noted that a P2 N-methyl glycine (secondary amine) was recognized as a DPP-IV substrate, suggesting -HNCH2CO-Pro- as a viable template for inhibitor design. As a result, tandem resinsolution parallel synthesis was carried out to investigate the activity of more complex N-substituted glycines coupled to an electrophilic (2S)-cyanopyrrolidine at P1. Resin-bound N-bromoacetylproline, 5, prepared in three steps from Rink Tandem solid/solution-phase synthesis of dipeptidyl peptidase IV (DPP-IV) inhibitors:



9: clinical candidate IC₅₀ = 7.0 nM (human plasma DPP-IV) >10,000-fold selective vs DPP-II (bovine kidney) and post-proline-cleaving enzyme (human PPCE); 75% *F* (rat, monkey); t_{1/2} of 100 mg oral dose in humans = 0.85 h with >80% inhibition of plasma DPP-IV for 4 h

Figure 1. Nitrile library yielding dipeptidyl peptidase-IV (serine protease) inhibitors.³⁶⁷

resin, was reacted with some 200 amines and cleaved to yield N-substituted Gly-Pro-NH₂ **6**. Library 1.18 (**3**) was obtained upon the solution-phase dehydration of **6** with trifluoroacetic anhydride (TFAA) followed by treatment with NH₃ in MeOH (**6** \rightarrow **7** \rightarrow **3**). One of the more potent human plasma DPP-



Figure 2. Nitrile library yielding cathepsin S (cysteine protease) inhibitors.³⁷⁴

IV inhibitors found in the library was the 5-nitro-aminopyridinyl glycine derivative 8: $IC_{50} = 8.7$ nM. In a followup medicinal chemistry effort, the potentially toxic 5-nitro group was replaced with a 5-cyano group in the pyridine ring to give orally active 9, the clinical candidate: $IC_{50} = 7$ nM possessing >10 000-fold selective against closely related proteases, including post-proline-cleaving enzyme (PPCE) and kidney DPP-II. In vivo evaluation of 9 in rats and humans definitively linked DPP-IV inhibition to sustained GLP-1 levels and an improvement in oral glucose tolerance.

Independently, Ward and co-workers at Boehringer Ingelheim developed a tandem resin-solution parallel synthesis of dipeptide-based nitriles as inhibitors of the cysteine protease, cathepsin S.³⁷⁴ This proteolytic enzyme is intimately involved in immune modulation by processing proteins (final processing step) which are presented as antigenic peptides on the surface of cells. Attenuation of self-antigen presentation via cathepsin S inhibition is of particular therapeutic interest for potential autoimmune disease intervention. Nitrile inhibitors were prepared by three different protocols, including one solid-phase protocol (illustrated in Figure 2) intended to define the preferred P1 specificity requirement for the enzyme. Fmoc-protected amino acids were coupled to piperidine-treated Sieber resin 10, affoding 11 after protecting group removal. Resin 11 was capped with the morpholino urea derivative of leucine 12 and cleaved from resin yielding amides 13. Primary amides 13, in turn, were converted to peptide nitriles 14 (library 1.21) using cyanuric chloride as the dehydrating reagent. Dissociation constants (K_d) increased with extended length of the P1 side chain $(15 \rightarrow 16 \rightarrow 17)$, with the benzyl-protected serine inhibitor 17 possessing a $K_{\rm d}$ of 300 pM. An X-ray crystal structure of the inhibitor 17-enzyme complex confirmed for the first time the putative covalent interaction between the active-site cysteine residue and the nitrile carbon atom, that is, thioimidate bond formation. The dipeptide nitriles prevented processing of the p10 invariant chain fragment (a cathepsin S substrate) in a cellular assay.

Fungal Type-1 Protein Geranylgeranyltransferase (GGTase-1) Inhibitors.^{346.} The dithiothreitol 18, a potent inhibitor of Candida albicans type I protein geranylgeranyltransferase (GGTase-1), was discovered through random screening at Banyu (Figure 3).³⁴⁶ GGTase-1 is an enzyme indirectly responsible for essential fungal cell wall biosynthesis via geranylgeranylation of certain G-protein regulatory components of (1,3)- β -D-glucan synthase. GGTase-1 is an α/β heterodimer containing both Mg²⁺ and Zn²⁺ in its active site, and the high affinity of 18 was attributed to the metal chelating properties of the dithiothreitol (DTT) mioety present in 18. Despite potent action against the enzyme, inhibitor 18 displayed weak antifungal activity (MIC > 120 μ g/mL). The DTT group was targeted as the barrier to whole cell activity. In an attempt to establish antifungal activity, libraries were designed to remove the DTT liability. For the first iteration, DTT esters were simplified to a single chelating thiol group, furnishing library 19. Subsequent screening of 19 afforded a new monothiol-containing inhibitor 20 with an IC₅₀ = 240 nM, albeit \sim 10 times less potent that the starting lead 18. Hoping to restore in vitro binding affinity, a second library design/synthesis iteration was carried out in which 3-mercaptopropanol was linked to trityl resin and acylated with a broad array of amino acids (library **21**; 70 members). This led to inhibitor **22** (IC₅₀ = 7 nM). The naphthylalanine and 1-indane moieties in 22 were essential for high enzyme affinity. Because of the researchers' dissatisfaction with ester thiol chelate, a third library iteration was carried out wherein the 1-indanyl-naphthylalanine groups were kept constant while the 3-mercaptopropanoate was replace by amino acids with side chain functionality capable of metal chelation. From the corresponding library 23 (266 members), D-histidine analogue 24 was identified as a marginally active GGTase-1 inhibitor. The descarboxamide analogue 25 was subsequently prepared as a discrete compound, leading to a 10-fold increase in activity (IC₅₀ \sim



Figure 3. Evolution of imidazole-based geranylgeranyltransferase (GGTase) inhibitors.³⁴⁶

3000 nM (24) versus $IC_{50} = 350$ (25)). This prompted the final design iteration linking histamine to a solid support and generating some 1200 compounds, as per library 2.18 (26). Screening this library revealed D-3,4-dichlorophenylalanine as a surrogate for the D-naphthylalanine, provided the amino acid was capped with a 9-xanthenyl moiety, for example, 27. The non-thiol-containing inhibitor 27 possessed an IC_{50} of 10 nM, comparable to the starting DTT-containing lead **18** (IC₅₀ = 17 nM); however, **27** also lacked antifungal activity against C. albicans. The poor in vivo activity was rationalized on the basis of new information that C. albicans can sustain growth in the absence of GGTase-1 and, hence, was not a function of inhibitor structure. Although compound 27 no longer contains a thiol group, it was arrived at through a rather circuitous library cascade. Perhaps it would have been more efficient to simply recognize imidazole as a metalchelating ligand and attach it (histamine) directly to the resin for derivatization negating the need for libraries **19**, **21**, and **23**. No in vitro activity was reported for the obvious analogue of **18** in which DTT-type chelate was exchanged for the histamine-type chelate.

KDR (VEGFR-2) Receptor Tyrosine Kinase Inhibitors. As part of an oncology program directed toward the discovery of antiangiogenesis agents, Merck conducted a high-throughput screening campaign against KDR kinase and identified pyrazolo[1,5-*a*]pyrimidine **28** ($IC_{50} = 224$ nM) as a novel small molecule lead (Figure 4).¹³⁰ KDR is a receptor tyrosine kinase mediating mitogenic signaling of vascular endothelial factor 2 (VEGF-2). VEGF is one of several growth factors that tumors express under hypoxic stress, ultimately triggering an angiogenic response. Chemistry was charged with the goal of improving the activity of **28**. To



Figure 4. Merck's KDR kinase inhibitors.¹³⁰

this end, a three-step solution-phase synthesis of this compound class was developed. The key step was the condensation of 3-amino-4-arylpyrazoles 31 (prepared in two steps from any lacetonitriles and hydrazine: $29 \rightarrow 30 \rightarrow 31$) with commercially available 2-arylmalonates 32. Heating 31 and 32 in the presence of acetic acid in ethanol and cooling yielded the crystalline 3,6-diarylpyrazolo[1,5-a]pyrimidines 33 (library 2.1). SAR obtained from library screening indicated that substitution at the para position of the 6-aryl ring enhanced inhibitory activity ($28 \rightarrow 34$; ~10-fold increase in affinity). Further analogue synthesis afforded 35-37. In general, these inhibitors were relatively nonselective for KDR versus PDGFR β , FLT-1, and FLT-4 kinases, but moderately selective versus FGFR-1 and Src kinases. Compound 36 inhibited VEGF-stimulated mitogenesis in human umbilical vein endothelial cells with an IC50 comparable to its in vitro KDR IC₅₀.

KDR inhibitors were also identified from a 23-member library of aminothiazoles **45** (library 2.2) synthesized by Waldmann and co-workers, who employed a traceless hydrazine linker strategy (Figure 5).³³⁸ The synthesis initiated with coupling (DIC, HOBt) commercially available nitrophenylhydrazines **39** (two inputs: **46** and **47**) to customprepared adipicate-functionalized resin **38**. Bis(Fmoc) protection of the hydrazide nitrogen atom (**40** \rightarrow **41**) preceded

nitro group reduction (SnCl₂·H₂O, DMF). Subsequent reductive amination of 41 proved difficult, but proceeded quantitatively with 10 equiv of R^2 CHO (five inputs: 48–52) and NaCNBH₃ in THF/AcOH (100:1), with imine formation and reduction occurring in sequential reaction steps. Treatment of the corresponding resin with commercially available Fmoc thioisocyanate followed by global Fmoc deprotection furnished the penultimate intermediate resin 42. This resin reacted cleanly with α -bromoaryl ketones 43 (three inputs: 53-55) under Hantzsch-type cyclization conditions and gave resin-bound thiazoles 44. Traceless cleavage $(44 \rightarrow 45)$ was accomplished by treatment with a catalytic amount of Cu- $(OAc)_2$ in *n*-propylamine and purging with O_2 to reoxidize Cu⁺ generated during the oxidation of the linker group. Copper salts were separated (99.9%) from the product by scavenging with cooper-chelating polyamine resin or through solid-phase extraction. Product purity (81-99%) was sufficient for biological screening. Because it was recognized that 2-aminothiazole derivatives are known inhibitors of cyclin-dependent kinases (CDKs), the library was initially screened against CDK-2 and CDK-4. However, none of the compounds proved active in the assays. Upon considering the high homology of the ATP-binding domain within the class of protein kinases, library 45 was evaluated against several receptor tyrosine kinases. These included epidermal growth factor receptor (EGFR; ErbB-1), ErbB-2 (Her-2/Neu), insulin-like growth factor 1 receptor (IGF1R), fibroblast growth factor receptor 1 (FGFR1), vascular endothelial growth factor receptors 2 and 3 (VEGFR-2 (KDR) and -3), and Tie-2. Inhibitory activity was observed for seven compounds against Tie-2, KDR, VEGFR-3, and FGFR-1. Inhibitor 56 was active against both KDR (IC₅₀ = 7.4 μ M) and Tie-2 (IC₅₀ = 9.8 μ M), while compound **57** (IC₅₀ = 12 μ M) was 3.5-fold selective for KDR versus VEGFR-3 and inactive against Tie-2.

Dynamic Combinatorial Library of Neuraminidase Inhibitors. Researchers at Therascope AG described an elegant application of dynamic combinatorial chemistry to the discovery of inhibitors of the vial enzyme neuraminidase (Figure 6).¹⁶⁴ Tamiflu **58a** is a marketed antiviral that targets influenza A virus neuraminidase. The viral surface enzyme



catalyzes the cleavage of sialic acid residues terminally linked to glycoproteins and glycolipids and plays an important role in the propagation of the virus. SAR and X-ray crystallographic studies have shown that the carboxylate, amino, and acetamido groups are salient binding elements and that there exists a hydrophobic pocket adjacent to the active site, partly occupied by the alkyl ether of **58a**. A dynamic combinatorial library of potential inhibitors was designed on



Figure 5. Aminothiozole-based dual active KDR and Tie-2 kinase inhibitors.³³⁸

the basis of diamine scaffold **58b**, in which the amine functionality was subject to reductive amination with lipophilic aldehydes. This chemistry would give rise to potential inhibitors that may engage the hydrophobic pocket, mimicking the alkyl ether in **58a**. In a control experiment, mixing **58b** with an aldehyde produced an equilibrium mixture of hemiaminal species **59** and **60**, as evidenced by NMR. Upon treatment with tetrabutylammoniumcyanoborohydride, the corresponding mono-, di-, tri- and tetrareductive amination products **63–66** were all detected by HPLC/MS. When the

reaction was carried out with a mixture of aldehydes in the presence of enzyme, target-biased reequilibration occurred toward species **59**, and amplification of but a few monoreductive amination products were detected. Resynthesis of the amplified products gave **67–69** with micromolar activity against the enzyme. Repeating this chemistry with BSA in place of enzyme did not produce any amplified products, indicating that legitimate target-driven selection/pressure was operating. In further experiments, scaffold **70** was mixed with a set of 20 aldehydes, producing up to 40 000 virtual



Figure 6. Neuraminidase inhibitors from a dynamic combinatorial library 2.27.¹⁶⁴

compounds in a single pot from which only a few amplified products were obtained, for example, **71** and **72**. Targetdriven self-organizing virtual libraries are envisaged for other systems, provided the target may be obtained in sufficient quantity and reaction conditions developed in aqueous buffer to allow transient equilibrium components to be formed in only trace amounts.

Substituted Indoles as Privileged GPCR Ligands. A mixture synthesis of 128 000-member library **81** (library 3.20) of 3-(amidoalkyl)- and 3-(aminoalkyl)-2-arylindoles was reported by Merck (Figure 7).³⁸⁰ The rationale for its construction was derived from the common occurrence of

indole derivatives in biologically active agents. The indole ring system is regarded as a privileged motif for GPCRs.⁴¹⁵ The design of library **81** was conceptually straightforward: (a) tether a basic amine to the 3-position of the indole ring; (b) introduce small functional group substituents in the 4-, 5-, 6-, and 7-positions; and (c) substitute the 2-position with a variety of aryls. Library construction was initiated by coupling arylketo acids **74** to Kenner safety catch linker **73** (Figure 7a). Condensation of resin **75** with aryl hydrazines **76** under conditions previously reported by the Merck group (ZnCl₂, AcOH) furnished the fully substituted indole scaffold **77**. The resin–SO₂–NH–CO– linkage was alkylated using



<u>Selected Z (-NR¹R²) pools and biological activity:</u> (Numbers in columns are % inhibition values at the given screening concentration)



Figure 7. Mixture synthesis and screening results of substituted indoles (library 3.2) as privileged GPCR pharmacophores.³⁸⁰

a Mitsunobu protocol (pentafluorobenzyl alcohol, PPh_3 , DIAD, THF) and treated with amine nucleophile to afford

the 3-(amidoalkyl)-2-arylindoles **79**. The corresponding 3-(aminoalkyl)-2-arylindoles **80** were prepared via post resin



- goal of finding lipophilic replacement for (A) in lead 90 achieved
 - no PK data given

Figure 8. Roche's VCAM/VLA4 antagonists.73

amide reduction with borane-dimethyl sulfide complex (BMS, 50 °C) and azeotropic removal ($3 \times$ with HCl/MeOH 50 °C) of borane byproducts. After each combinatorial step and prior to pooling, a portion of each individual resin was archived for later deconvolution as per standard combinatorial mixture technique. In total, some 128 000 compounds were prepared as 320 pools of 400 compounds.

To exemplify the power of mixture synthesis and the privileged nature of the indole scaffold, the compound pools were screened for biological activity at a single concentration (from 0.1 μ M up to 20 μ M screening concentration) across a panel of 16 GPCRs, including many receptor subtypes. Activity was observed against every receptor, and selectivity was highly dependent on the pattern of indole substitution. The most active pools from seven representative screening

hits were deconvoluted, yielding potent and highly selective ligands (Figure 7b). For example, **88** had an IC₅₀ of 0.6 nM against the neurokinin 1 receptor (NK₁), displaying > 10 000fold selectivity versus NK₂ and NK₃. Compound **86**, possessing an IC₅₀ of 0.8 nM for neuropeptide Y5 (NPY₅), was similarly > 10 000-fold selective versus NPY₁. Impressive, too, were the active compounds deconvoluted against the 5-HT receptors: for example, indole **84** has an IC₅₀ of 10 nM against 5-HT_{2a} and was 6-fold selective over 5-HT_{2c}; 60-fold selective over 5-HT₆; and >100-fold selective versus 5HT_{1a}, 5HT_{5a}, and 5-HT₇.

VLA-4 ($\alpha_4\beta_1$ Integrin) Antagonists. The very late antigen 4 (VLA-4) is a heterodimeric integrin ($\alpha_4\beta_1$) expressed on many lymphocytes, including T-lymphocytes and eosinophils. VLA-4 mediates cell adhesion, infiltration and activa-



Figure 9. Merck's VCAM/VLA4 antagonists.²²⁸

tion of these cells when it encounters various receptors, such as vascular cell adhesion molecule-1 (VCAM-1), at sites of inflammation. VLA-4 antagonists are expected to be of significant therapeutic benefit for the treatment of inflammatory diseases. Three companies, Merck,²²⁸ Roche,⁷³ and Celltech,²⁹⁹ reported on the synthesis and evaluation of focused libraries for identification of VCAM/VLA-4 antagonists with improved pharmacokinetic properties (lowered clearance). Each company started with structurally similar antagonists displaying high potency and integrin selectivity (Figures 8-10). The lead compounds, 90, 101, and 115, share a derivatized phenylalanine motif as the key pharmacophore. Each company arrived at their lead through screening or structure-based design exercises and had been pursuing VLA-4 antagonists for several years. Finally, leads 90, 101, and 115 are rapidly cleared from the blood upon iv administration, and each research group independently

hypothesized that the rapid clearance of the compounds may due to the N-acylated functionality (dipeptide-like character); hence, focused libraries were constructed modifying this region of the lead.

The approaches each company took for library synthesis were quite similar. The phenylalanine carboxylate was the functional group handle for solid-phase attachment. In the case of Roche (Figure 8), Wang resin was esterified with the orthogonally protected phenylalanine derivative **91**, and through a series of selective deprotections, they arrived at the 42-member library **94** (library 4.9). The Roche group found *N*-benzylpyroglutamide could be substituted by the less peptidic but more lipophilic 2,6-disubstituted benzamide (e.g., **99**; IC₅₀ = 1.2 nM) and α , α -spiroalkylphenyacetamide (e.g., **100**; IC₅₀ = 1.7 nM) moieties. These moieties bear remarkable similarity to the amide replacements that Merck found (see below). No pharmacokinetic data was given, but presumably, these agents were subject to rapid clearance.

From earlier studies, Merck observed that the 2-methoxyphenyl group in lead 101 was quite optimal for integrin antagonism (Figure 9). The biphenyl moiety substantially reduces the peptide character relative to the Roche series (4-amido linkage in 90 removed) and is arguably a better starting point for pharmacokinetic optimization. In an effort to reduce molecular weight and improve clearance, library 4.8 (102) was prepared. Wang resin was coupled with the Fmoc-protected 4-(2-methoxyphenyl)phenylalanine 103. Following Fmoc deprotection $(103 \rightarrow 104)$ and acylation (104) \rightarrow 102), the ~100 member library 102 was obtained. Surrogates for the sulfonated proline residue included 2,6disubstituted phenylacetamides (e.g., 106; IC₅₀ = 8 nM), heteroarylacetamides (e.g., 109: IC₅₀ = 12 nM), and α , α spiroalkylphenyacetamides (e.g., 111; $IC_{50} = 3 nM$). Hybrid structures 113 and 114 (IC₅₀ = 0.5 nM, each), combining elements of both heteroaryl and α, α -spiroalkyl groups and synthesized as discrete compounds based on library SAR, were the most potent VLA-4/VCAM1 antagonists. Despite a reduction in molecular weight, these agents still displayed unfavorably high clearance.

Celltech took a slightly different tack from Merck and Roche and sought an isosteric replacement for the thiazolidinecarboxamide in lead 115 (Figure 10). In particular, the researchers were interested in incorporating the amino acid isostere, 3,4-diamino-3-cyclobutene-1,2-dione. Library construction began with acylation of Wang resin with Fmocprotected 4-nitrophenylalanine 117. Nitro reduction proceeded smoothly to yield the corresponding resin-bound 4-aminophenylalanine, which was acylated with 3,5-dichloropyridine-4-carboxylic acid chloride 118 (117 \rightarrow 119). After Fmoc deprotection, resin-bound amine was reacted with 3,4dimethoxycyclobuten-1,2-dione 120 to give the penultimate intermediate 121. Treatment of 121 with primary and secondary amines followed by cleavage with 60% TFA/ DCM, and HPLC purification furnished a series of libraries, 116a-c, defined by the amine nucleophile (primary un-



Figure 10. Celltech's VCAM/VLA4 antagonists.²⁹⁹

branched, primary branched, or secondary amine). In general, the squaric acid derivatives displayed good potency, again demonstrating their utility as amino acid surrogates. Of special interest was *N*-propyl analogue **123** (IC₅₀ = 1.5 nM (protein-based assay); IC₅₀ = 120 nM (whole cell assay), which showed high potency and reduced iv clearance (0.1 mL/min/kg (rat)). Interestingly, Celltech prepared a squaric acid analogue of Merck's 2-methoxyphenylphenylalaninecontaining antagonist **101** and was found to have diminished potency by a factor of ~10 (whole cell assay), suggesting a unique SAR for the squaric acid series.

Diversity-Oriented Synthesis (DOS). There were several architecturally sophisticated DOS libraries originating from the laboratories of Stuart Schreiber at the Harvard Institute of Chemistry and Cell Biology.^{213,217,218,336} This group has been developing encoded split synthesis of stereochemically complex and diverse small molecules. The libraries are not necessarily intended to be druglike in terms of overall physiochemical properties or yield drug leads per se, but rather, they are natural product-inspired designed to explore cellular and organismal pathways vis a vis chemical genetics. Chemistries are typically carried out on high-capacity mac-

robeads (500 μ m; capacity of 100 nmol per bead), providing a sufficient amount of compound after cleavage and formulation as a "one bead/one stock" solution for screening in many assays. A recurring linker theme is the immobilization of functionalized alcohol- or phenol-containing building blocks by way of a silyl ether bond (macrobead-alkylsilyl triflate). After single bead arraying, release of library compounds is achieved by exposure to HF-pyridine. Excess reagent is removed by the addition of TMSOMe, followed by concentration under reduced pressure. All library members therefore possess a hydroxyl group in their structure. Compound identity and purity is ascertained by LCMS analysis of a small portion of cleaved material, or in those cases in which libraries are encoded, GC analysis of the unique set of electrophoric tags revealing the bead's chemical reaction history. Encoded split synthesis library 136 of 29 400 discrete, polycyclic compounds relied on differential dienophile reactivity with resin bound trienes to achieve skeletal diversity (Figure 11).²¹⁸ Some 14 X-ray crystal structures of library compounds were determined as a visual aid to reveal skeletal diversity. Ferrier and Pauson-Khand reactions on a glycal template led to a 2500-member encoded library 151



Figure 11. Diversity-oriented synthesis of polycyclic compounds (tags not shown for clarity).²¹⁸

of tricyclics (Figure 12).²¹³ The efficient oxidation of resinbound organocuprates was exploited in the atropdiastereoselective synthesis of biaryl-containing medium rings (9-, 10-, and 11-membered ring; library 4.20 (163-165; Figure 13).³³⁶ Electrophoric tags were incompatible with its library chemistry (metalation of the polychlorinated tags occurred during biaryl bond formation), making encoded split-pool synthesis impossible. In this instance, care was taken to avoid building block constitutional isomers, (S)- and (R)-amino alcohols were processed in parallel libraries, and bead pooling was eliminated in the final synthetic step to ensure that compound identification could be secured by electrospray mass spectrometry. Biological evaluation of DOS-type libraries has yielded numerous novel small molecules having specific and potent protein-binding and cellular activities, representing potential probes for chemical genetic studies.⁴¹⁶

Resin-bound 2*H*-pyran-3(6*H*)-one intermediates **171** were conceived as a multireactive core molecule from which a variety of pharmacophoric frameworks could be created (Figure 14).¹⁰⁰ In the work described by Couladouros and Strongilos, a series of α -hydroxyfurans **166–169** with an oxidation-sensitive (DDQ) linker was synthesized in solution and then immobilized to give resin **170**. The derivatized resin was then treated with NBS, THF/H₂O (4:1) at 0 \rightarrow 25 °C for 1 h. Pyranones **171** so obtained were subjected to a host of chemistries, leading to both skeletal rearrangements and function group interconversions $(171 \rightarrow 172 \rightarrow 182)$. Structurally diverse mono, di-, and tricyclic heterocycles were all obtained from 171.

Solid-phase synthesis directed toward creating diverse collections of heterocycles has been reported. Strategies deployed to date require synthetic sequences to converge to a common intermediate or set of intermediates, which then diverge into different heterocyclic products. Purandare (Bristol-Myers Squibb) converted o-fluoronitrobenzoic acid into resin-bound aniline 187 and proceeded to construct six heterocyclic ring systems (188–193; Figure 15).³⁰⁴ Huang (Shanghai Institute) converted Merrifield resin to a polymerbound variant of Meldrum's acid 199 (Figure 16).¹⁷³ Condensation of 199 with triethylorthoformate generated the corresponding aminomethylene cyclic malonic acid esters 200. Reaction of 200 with either an aniline, urea, or thiourea then intracyclative cleavage (thermolysis: 220-240 °C, 20 min) furnished quinolones 201, pyrimidobenzothiazolones 202, thiazolopyrimidinones 203, uracils 204, and thiouracils 205, respectively.

Solid-Phase Synthesis Using Katritsky's Benzotriazole Chemistry. Nearly two decades ago, Katritzky began an odyssey exploring the utility of benzotriazole (Bt) in carboand heterocyclic synthesis and has published extensively in



Figure 12. Diversity-oriented synthesis of tricyclic compounds.²¹³

this area. Bt-based chemistry is finding its way into solidphase synthesis. There were eight Bt-related publications appearing from four independent research groups in 2002.

Katritzky disclosed the solid-phase synthesis of 4,6-diaryl substituted and 3,4,6-trisubstituted pyrid-2-ones **213** by reaction of resin-bound chalcones **211** with 2-(benzotriazol-1-yl)- and 2-alkyl-2-(bentriazol-1-yl)acetamides **212** (Figure 17A).¹⁹³ Resin-bound chalcones **211** were prepared by linking 4-hydroxyacetophenone **208** to Wang resin **207** by a modified Mitsunobu protocol (DIAD, PPh₃, NMM, 25 °C, 48 h) followed by condensation with aryl aldehydes (MeONa, THF, 25 °C, 24 h). Pyridones **213** were obtained in excellent yield and purity upon reaction of **211** with 2-(benzotriazol-1-yl)acetamides (20 equiv) in the presence of NaOH (40

equiv) in EtOH/THF (1:5) at 70 °C for 24 h, after standard TFA cleavage conditions.

Katritzky reported the condensation of amine resins **215** with di(benzotriazol-1-yl)methanimine **216** (25 °C, 7 h) to give Bt-containing intermediates **217** (Figure 17B).¹⁹⁴ Treatment of **217** with thiols or secondary amine activated with EtMgBr lead to *S*-arylisothioureas **218** and guanidines **219**, respectively.

Finally, Katritzky demonstrated that *N*-acylbenzotriazoles **220** can be employed in the solid-phase preparation of amides (Figure 17C).¹⁹⁴ Wide structural variation in the R2 group was geneated by reaction of *N*-methansulfonylbenzotriazoles with a carboxylic acid in the presence of base. Resin-bound amines **215** were acylated under neutral conditions upon



Figure 13. Diversity-oriented synthesis of biaryl-containing medium rings.³³⁶

treatment with **220** in refluxing THF for 48 h. Resin cleavage provided the correspoding amides **221**.

The versatility of benzotriazole-1-carboximidamide-type intermediates **222** was further demonstrated by Makara and co-workers in a regiocontrolled synthesis of 3-alkylamino-1,2,4-triazoles **224** and 1,2,4-oxadiazoles **225** (Figure 18).^{248,249} Inspired by Katritzky's original solution synthesis, acylation of **222** with alkyl and aryl acid chlorides generated immobilized *N*-acyl-Bt-1-carboximidamides **223**. Reaction of **223** with N-substituted hydrazine or hydroxylamine with DBU as the optimal base in THF proceeded at 50 °C to give the respective heterocycles. DBU was essential to ensure efficient cyclization and minimal byproduct formation.

A facile route to 1,2,5-trisubstituted 4-imidazolidinones was reported by Houghten (Figure 19).³¹⁹ MBHA resinbound amino acids **226** were prepared via classical amino acid coupling/Fmoc deprotection/reductive amination methodology. When resin **226** was refluxed with an aromatic, aliphatic, or heterocyclic aldehyde (10 equiv) and benzo-triazole (10 equiv) in benzene for 16 h, the reactive *N*-[1-

benzotriazol-1-yl)alkyl] species 227 was produced, which suffered spontaneous intramolecular cyclization via nucelophilic substituion of the Bt group with the amidic nitrogen $(226 \rightarrow 227 \rightarrow 228)$. Treatment of resin 228 with HF released compounds 229. The reaction is nonstereospecific and produces diastereomers in ratios that vary depending on the ring substitutents. The high-yielding reaction conditions and range of building blocks lends itself to the production of large, diverse collections of imidazolidinones as single compounds or in mixture-based synthesis. This group also reported the application of this chemsitry to the synthesis of peptidomimetic-based 4-imidazolidinones.³¹⁶

Researchers from the University of Amsterdam were interested in creating libraries of 2-substituted piperidines **232** as potentially bioactive compounds (Figure 20).³⁶⁶ An efficient solid-phase synthesis was developed starting from carbamate-tethered δ -amino acetals **230**. The key reaction was an intramolcular cyclization to form a transient *N*-acyliminium ion **231** that could be trapped with suitable C-nucleophiles. Lewis acidic conditions previously successful



Figure 14. Diverse pharmacophores from resin-bound 2H-pyran-3(6H)-ones.¹⁰⁰

in a high-yielding solid-phase preparation of 2-substituted pyrrolidines from γ -amino acetals failed with the homologous piperidine analogues. For example, although the reaction with **233** and allylsilane gave the expected 2-piperidine, it was contaminated with a linear side product resulting from competing intermolecular attack of the allylsilane onto the oxycarbenium ion instead of the intramolecular attack by the carbamate nitrogen atom. This side reaction was not observed in the pyrrolidine case. Several attempts were made to suppress the reaction, but to no avail. Model system **233** was then studied in solution. Treatment of **233** with BF₃· Et₂O led to a mixture of **235** and the corresponding linear product **236**. Exposure of **233** to catalytic *p*-TSA in DCM

or DCM/EtOH produced rather unstable intermediates **237** and **238**. However, the reaction of **233** with catalytic *p*-TSA in DCM for 15 min at room temperature followed by the addition of BtH **206** led to quantitative conversion to the 2-Bt-substituted piperidine **239**. Bt derivative **239** was stable to silica gel column chromatography and could be fully characterized. Treatment of **239** with a Lewis acid and allyITMS furnished **235** in 93% isolated yield. These reaction conditions were then applied to resin-bound acetal **230**. Complete conversion to the putative ring-closed intermediate **240** was established by IR (disappearance of the NH signal). Subsequent treatment of **240** with a range of C-nucelophiles and cleavage with 1 M NaOMe in THF/MeOH (2:1) gave



Figure 15. Divergent heterocyclic synthesis from a common resin-bound intermediate.³⁰⁴



Figure 16. Diverse heterocycles from resin-bound Meldrum's acid.¹⁷³

2-substituted piperidines **232** in good yield and purity and in high diastereoselectivity.

Amine Synthesis. Amines and derivatives thereof are ubiquitous in drug substances, and combinatorial methods are continually sought to generate this salient functional group. Examples of new protocols for the preparation of secondary and tertiary amines and the N-arylation of primary and secondary amines were reported in 2002.

Pelletier and co-workers at Wyeth described a novel, tandem three-phase reductive amination procedure for the preparation of secondary amines from primary amines with minimal side product formation (Figure 21).²⁹¹ Typical side products obtained from classical solution or solid-phase reductive amination conditions include (a) alcohol from reduction of the carbonyl group, (b) starting amine from incomplete reaction, and (c) products of over alkylation, all of which may diminish yield and compromise product purity. In Pelletier's multiphase reaction, resin-bound aldehyde 242 in its resin-bound imine form 244 undergoes a transimination/ cleavage equilibrium with a primary amine 245 (0.5 equiv) to produce a solution-phase imine 246. In the presence of polymer-bound borohydride (2 equiv), the solution imine 246 is reduced to the corresponding secondary amine. After filtration, excess starting resin 243, resin-bound imine 244, and reducing reagent are removed, leaving essentially pure product. Because there is no excess aldehyde present and the product (secondary amine) does not participate in the initial phase switch, over alkylation does not occur; neither is any alcohol from RCHO reduction observed. The methodology was applied to a library of β -adrenergic receptor ligands; compounds 249-251 are representative members. It will be of interest to see whether other research groups



Figure 17. Application of Katritzky's benzotriazole to solid-phase synthesis.^{193–195}

who attempt this simple, yet elegant, chemistry will find equally satisfying results.

A novel and versatile safety catch linkage strategy was developed by Schultz to generate libraries of functionalized *N*,*N*-dimethyltryptamines and β -carbolines (Figure 22).³⁸⁶ Tryptamine scaffolds **252** (prepared in three steps from commercially available indoles) were immobilized through



Figure 18. Benzotriazole-mediated triazole and oxadiazole synthesis.^{248,249}



Figure 19. Benzotriazole-mediated 4-imidazolidinone synthesis.³¹⁹

the Michael addition to vinyl sulfone resin 253, furnishing tryptamine-bound resin 254. In model studies, activation of the safety catch linker by exhaustive methylation (MeI, 15 equiv) and Hoffman elimination mediated by DIEA proceeded smoothly to furnish N,N-dimethyltryptamines (254 \rightarrow 255 \rightarrow 256). Alternatively, clean monomethylation to resin 257 was achieved upon brief exposure to a slight excess of MeI (2 equiv, 15 min). This resin 257 was sufficiently stable toward acylating reagents, Cu-catalyzed N-arylation and Suzuki coupling conditions permitting further scaffold diversity $(257 \rightarrow 258/259)$. Last, resin 254 was subjected to a Pictet-Spengler cyclization (257 \rightarrow 260). Premature cleavage of material from the solid support was not observed. Quaternary ammonium formation and Hoffman elimination then generated the *N*-methyl- β -carbolines **261**. Excellent purity and yields were obtained for all transformations.

Combs at DuPont Pharmaceuticals, now Bristol-Myers Squibb, reported additional details on an operationally simple and mild method for the N-arylation of resin-bound primary and secondary amines (Figure 23).⁹⁴ *N*-arylamines are widely found in biologically active agents and, hence, are attractive pharmacophores for library exploitation. Copper acetatepromoted C/N cross-coupling of arylboronic acids with

Synthetic strategy:



Figure 20. Benzotriazole-mediated piperidine synthesis.³⁶⁶

amines takes place via sequential addition of phenylboronic acid (4 equiv), anhydrous copper acetate (2 equiv), 4Å powdered molecular sieves (50 wt % of resin), and triethylamine (4 equiv) to the resin-bound amines (e.g., 262) in THF. The cross-coupling reaction proceeds for \sim 3 h, after which time the resin is drained and washed. This is repeated for a total of three cycles to maximize formation of product 263. Conversion to N-arylated products was on the order of 70-90% for para-substituted boronic acids, whereas lower conversions (35-50%) were observed for other substitution patterns. Overall yields were moderate to good after HPLC purification of the products (20-75%). One of the interesting features of this cross-coupling chemistry is that primary amines that lack an α -substituent are efficiently converted to the N,N-diaryl products $(264 \rightarrow 265)$ in high yield with <10% mono-N-aryl product fromation. Introduction of α -substitution, a small methyl group, for example, completely changes the product profile, and only mono-N-arylamines $(266 \rightarrow 267)$ are formed without any trace of the corre-



Figure 21. Tandem three-phase reaction for secondary amine

sponding diarylamine. Sulfonamides and many heterocycles containing free N-H bonds afford good to excellent yields and purities of N-arylated products. In contrast, amides, carbamates, and ureas give no reaction.

synthesis.291

Fluorous-Based Reagents and Scavengers. Fluorous chemistry was pioneered by Curran⁴¹⁷ and has formed the basis of a relatively new startup company, Fluorous Technology, Inc. (FTI). The basic tenet underlying the technology is that molecules tagged in some fashion with a perfluoroalkyl group can be rapidly, selectively, and quantitatively separated from molecules without such tags. The unique physiochemical properties of the perflouroalkyl group override other functionality present in the molecule. Liquidliquid extraction (fluorohydrocarbon solvent) and chromatography on fluorous sorbants are employed to separate fluorous tagged/untagged compounds. The nascent technology is finding application in high-throughput synthesis, particularly in the parallel synthesis of discrete compounds. Combinatorial mixture synthesis is also possible using perfluoroalkyl groups of varying chain lengths.⁴⁰⁹ Fluorous tags are inert to most chemical reaction conditions and offer a convenient handle for rapid separation, and chemistries can

N,N-dimethyltryptamine synthesis



Figure 22. Safety-catch linkage strategy for the synthesis of N,N-dimethyltryptamines and β -carbolines.³⁸⁶



a: 1) Cu(OAc)₂ (2 equiv), ArB(OH)₂ (4 equiv), TEA (4 equiv), THF, 4 Å sieves, 3 cycles total; 2) TFA/DCM (1:1)

| Figure 23. | N-Arylation | of primary | and secondary | amines o | n solid |
|------------|-------------|------------|---------------|----------|---------|
| support.94 | | | | | |

be performed in solution and are easily scaleable. A growing collection of fluorous reagents, protecting groups, miscellaneous tags, and assorted fluorous sorbants, for example, fluorous SPE cartridges (Fluoro*Flash* SPE), for purification are commercially available.

The Technology Enabled Synthesis Group at Merck has been actively developing fluorous-tethered reagents and scavengers to facilitate parallel solution synthesis. A novel fluorous variant of triphenylphosphine in combination with FluoroFlash SPE was used in a Staudinger protocol for azide to amine reduction.²³⁵ The reduction reaction was monitored by ³¹P NMR, and reaction times of only 4 h were necessary, in contrast to >72 h required for resin-bound triphenylphosphine. A series of fluorous-tethered quenching reagents 268-273 were prepared and successfully utilized in several reaction classes to scavenge excess amine, aniline, alkoxide, and epoxide functionality from solution. For example, in the ureidation of benzylamine with excess phenethylisocyanate, Lindsley²³³ demonstrated that excess phenethylisocyanate was efficiently scavenged by C_7F_{15} -tagged amine 268, affording urea 275 in 93% yield and >98% purity after FluoroFlash SPE chromatography (Figure 24).

The commercially availability of fluorous-tagged acid chlorides and a great number of diamines led to the design and synthesis of scavengers 277-282 by the Merck group.²³⁴ Simple acylation chemistry offered a convenient source of

<u>Fluorous scavengers removed from reactions with</u> FluoroFlash[™] SPE alone:²³³



Fluorous-tethered bases **not** removed from reactions with FluoroFlash[™] SPE alone. **277-282** require an ion exchange precolumn due to polar functionality.²³⁴



Figure 24. Fluorous-tethered scavengers for solution-phase parallel synthesis.^{233,234}

these and potentially many other custom fluorous-tethered bases to complement existing resin-bound congeners and to capitalize on homogeneous solution-phase kinetics. However, during the initial evaluation of **280** in a sulfonylation reaction, $\sim 15\%$ of the fluorous base coeluted with sulfonamide product when the reaction was purified by Fluoro*Flash* SPE. It came as a great surprise to discover that none of the fluorous bases **277–282** were fully retained upon Fluoro*Flash* SPE. Even with a silica gel transfer precolumn in place, up to 60% of some for the fluorous amide scavengers passed though the fluorous SPE cartridge. This stands in contrast to current dogma that the chromatographic properties of the perflouroalkyl group dominate over other functionalities present in the molecule. Lindsley noted all previously



Figure 25. Ugi-type condensation using sulfonamide as amine input.⁶⁰

reported fluorous-tagged reagents have the perfluoroalkyl group linked in close proximity to the scavenging functionality by a lipophilic tether, whereas scavengers 277-282 contain a polar amide bond and position a basic nitrogen several atoms away from the perfluoro tag. Ultimately, a "mixed sorbent" solved the problem: crude reaction mixtures were first passed through an ion exchange column followed by fluorous SPE. In this way, products were isolated in high yield and desired purity (>95%, free of fluorous tags). In addition, a set of guidelines based on LCMS retention time was developed to predict a fluorous-tethered reagent's chromatographic behavior on Fluoro*Flash* SPE.

Multicomponent Condensations. Multicomponent condensations give rise to products with several points of diversity by bringing together three or more different building blocks in a single reaction step. The Ugi reaction, popularized in recent years by combinatorial chemists, utilizes four components, a carboxylic acid, an amine, an aldehyde, and an isocyanide, generating acylamino amides, which may be further transformed into a wide variety of acyclic and heterocylic products. A research group at Advance SynTech explored sulfonamides as surrogates for the usual amine building block in this reaction, potentially yielding α -sulfonylamino amide derivatives (Figure 25).60 The reaction was first examined in solution wherein a *p*-toluenesulfonamide was treated with 1 equiv each of acetic acid (HOAc), hydrocinnamaldehyde, and tert-butyl isocyanide in THF/ MeOH (1:1). After 3 days, a complex mixture formed with only trace amounts of the desired four-component condensation product detected. Despite the poor reactivity of the sulfonamide, the fact that trace quantity of desired material was formed spurred further investigation on solid-phase. Resin-bound sulfonamide 285 (from 284 and Rink resin, DIC/HOBt coupling) was treated in exactly the same way as in the solution experiment, except that 10 equiv of each





Figure 26. Hlasta's multicomponent 2-substituted azole synthesis on solid phase.¹⁰⁸

component was used. After 3 d, the resin was exposed to 20% TFA in DCM. Products so cleaved were examined by LC/MS, and the results again confirmed trace product **288**, but largely unreacted sulfonamide. The reaction was repeated, but this time at elevated temperature (60 °C, 24 h). Cleavage gave the desired Ugi product **288** contaminated with acylated materials **286a/b**. Decetylation occurred easily upon exposure to 40% aq MeNH₂/THF (1:1), affording complete conversion to α -sulfonamino amide **288** after resin cleavage. The reaction conditions were successfully applied to a broad range



Figure 27. Three-component condensation yielding substituted quinolines.¹⁷⁶

of aldehydes and isocyanides. The diversity of products could be further expanded by N-alkylation of **287** with alcohols under Mitsunobu conditions. Product yields for **289** were on the order of 40-93%, with purity ranging from 60 to 95% for 18 representative products.

Hlasta described a traceless solid-phase synthesis of 2-substituted azoles (Figure 26).¹⁰⁸ The three-component synthesis employs an azole (imidazole exemplified), aldehyde, and an amine. Imidazole 292 (3 equiv) is captured on solid phase by reaction with resin-bound carbamoyl chloride **291**. This generates an azolium ylide **293** which undergoes immediate condensation with an aldehyde (5 equiv, 24 h) present in the reaction medium. Intramolecular acyl transfer occurs, forming resin-bound 2-substituted imidazole (293 \rightarrow **294** \rightarrow **295**). Cleavage from resin using TFA in THF at reflux for 24 h in the presence of an amine, thiol, or alcohol nucleophile affords 296 via formation and trapping of an intermediate carbonium ion. In the absence of a nucleophile, alcohol **297** is produced. A 3 (heterocycle) \times 4 (aldehyde) \times 3 (nucleophile) library demonstrating the fidelity of the chemistry gave 34 out of 36 theoretical compounds, with 80% of the compounds possessing > 80% purity.

A novel three-component condensation protocol for the synthesis of quinoline derivatives was reported (Figure 27).¹⁷⁶ In a one-pot reaction, anilines, aromatic aldehydes, and a catalytic amount of CuCl in THF were stirred together for 15 min, followed by the addition of a terminal alkyne in THF (reflux, 10 h). Quinolines **310** were obtained in good yield after chromatography to remove benzylamine byproducts. Noteworthy is the mild reaction conditions and the use of propargyl alcohol and its acetate as an alkyne input. These inputs make provision for further derivatization chemistry.

Table 1. Chemical Libraries Targeting Proteases (Asterisk (*), Point of Attachment to Resin)



Activity: $k_{obs}/[I] = 28,735 \text{ M}^{-1}\text{s}^{-1}$



Y = amine

NH₂

Table 1. (Continued)

Library: 1.15

Name: Amino acid amide Size: >100 members Affiliation: Protherics [231] Note: *In silico* virtual library screening identified seed template which was elaborated using three iterations of library design.

Library: 1.16 Name: Indole carboxamide Size: *ca.* 50 members Affiliation: Aventis [254] Note: Part of a 138 compound library.

Library: 1.17 Name: β-strand mimetics Size: 48 members Affiliation: Fuchi, N.; *et al.* [131] Note: Constructed via 1,3-dipolar cycloaddition using resin-bound vinylsulfones.

Library: 1.18

Name: Pyrrolidine carbonitrile Size: *ca*. 200 members Affiliation: Novartis [367] Note: Corresponding primary amides (via Rink resin) were dehydrated to nitriles using trifluoroacetic acid anhydride.

Cysteine proteases

Library: 1.19 Name: Mercaptomethyl ketone Size: 20 members Affiliation: Huang, L.; *et al.* [172]

Library: 1.20 Name: Cyclic ketone Size: *ca.* 7 members Affiliation: Huang, L.; *et al.* [171]

Library: 1.21 Name: Dipeptide nitrile Size: Not defined Affiliation: Boehringer Ingelheim [374] Note: Multiple solution- and solid-phase libraries prepared to optimize R¹-R³.



NH₂

-AA-X-R

X= 0, NH

 H_2N

HN 🧖



NH



O

Enzyme: Factor Xa (human) Activity: K_i = 9 nM



Enzyme: Trypsin Activity: 65% inhibition at 100 nM

Enzyme: Dipeptidyl peptidase IV (human) Activity: IC_{50} = 22 nM



Enzyme: Cruzain Activity: K_i = 2.0 nM

0 Enzyme: Cruzain ö Activity: K_i = 16 nM

CN H Ô 0

Enzyme: Cathepsin S Activity: IC₅₀ = 5 nM



 $-NH_2$

ö Ń R1



Table 2. Chemical Libraries Targeting Nonproteolytic Enzymes (Asterisk (*), Point of Attachment to Resin)



`N∽ŃH

Enzyme: CDK4 Activity: IC_{50} = 260 nM, CDK4; IC_{50} = 7 nM, CDK2

`N-ŃН

Library: 2.4 Name: Imidazoguinoxaline Size: ca. 175 members Affiliation: BMS [74] Note: Library prepared using solution- and solid-phase synthesis.

Library: 2.5 Name: Pyrimidinylimidazole Size: 570 members Affiliation: Aventis [255] Note: Substrates linked to resin via Wang thiol linkage; S-oxidation then traceless cleavage with R³R⁴NH.



Library: 2.7 Name: Arylsulfonamide Size: ca. 500 members Affiliation: Reuveni, H.; et al. [312] Note: Multiple libraries optimizing ArSO₂-, diamine, and -CH₂R.

Library: 2.8 Name: Heterobiaryl Size: 324 members Affiliation: Novo Nordisk [267] Note: Two solution phase libraries.

Library: 2.9 Name: Heterobiaryl Size: >100 members Affiliation: Novo Nordisk [267] Note: Solution-phase follow-up library to library 2.8.

Library: 2.10 Name: Heterobiaryl Size: 96 members Affiliation: Novo Nordisk [267] Note: Solution-phase follow-up library to library 2.9.







Ar-SO₂-[diamine]-CH₂R

Enzyme: p56Lck Activity: IC₅₀ = 2 nM



Enyme: p38 MAP kinase Activity: IC₅₀ = 4 nM



Enzyme: Cyclin-dependent kinase 2 Activity: IC₅₀ = 3 nM

0,8

Enzyme: Protein kinase B/Akt Activity: $IC_{50} = 2 \mu M$





Enzyme: Glycogen synthase kinase-3 kinase-3 (GSK-3) Activity: IC₅₀ = 1.2 µM

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COOH

Activity: IC_{50} = 8.0 μ M



Enzyme: GSK-3 Activity: IC₅₀ = 0.4 μ M

Enzyme: GSK-3

Library: 2.11 Name: 3-Aminopyridazine Size: Not defined. Affiliation: Mirzoeva, S.; *et al.* [259] Note: Solution-phase parallel synthesis.

Phosphatases

Library: 2.12 Name: Difluoromethylene sulfonate Size: 24 members Affiliation: Leung, C.; *et al.* [226]

 CF_2

HO

R²HN

R₂

R

NH

78

ÌΝХ

Library: 2.13 Name: Difluoromethylene phosphonate Size: 20 members Affiliation: Hum, G.; *et al.* [175]

Library: 2.14 Name: Dysidiolide analog Size: 12 members Affiliation: Brohm, D.; *et al.* [52, 53] Note: Resin cleavage via olefin metathesis.

Dehydrogenases

Library: 2.15 Name: Pyrazole Size: *ca*. 280 members Affiliation: BMS [158] Note: Three focused libraries prepared.

Library: 2.16 Name: Substituted androsterone Size: 273 members Affiliation: Maltais, R.; *et al.* [251] Note: One lead finding library (168 members) and 2 follow-up libraries (56 and 49 members respectively).

Library: 2.17 Name: Substituted androsterone Size: 25 members Affiliation: Maltais, R.; *et al.* [251] Note: Follow-up to library 2.16.







Enzyme: Calmodulin-dependent kinase II Activity: IC_{50} = 9 μ M



Enzyme: Protein tyrosine phosphatase-1B (PTP1B) Activity: IC_{50} = 25 μM

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Enzyme: PTP1B Activity: K_i = 8 μM



Enzyme: cdc25c phosphatase Activity: IC₅₀ = 0.8 μ M



Enzyme: Dihydroorotate dehydrogenase (H. pylori) Activity: $K_i = 4 \text{ nM}$; >10,000x selective versus human enzyme.



Enzyme: Type 3 17 β -hydroxysteroid dehydrogenase Activity: IC₅₀ = 35 nM



Enzyme: Type 3 17 β -hydroxysteroid dehydrogenase Activity: IC₅₀ = 74 nM

 $N_{R^3} \longrightarrow K_{*R^1} \longrightarrow K_{*R^1}$

<u>Transferases</u>

Library: 2.18

Name: Histamine peptide Size: *ca.* 1200 members Affiliation: Banyu [346] Note: Initial lead was a dithiothreitol derivative. Sequential series of optimization libraries to arrive at potent inhibitor.

Library: 2.19 Name: Pepticinnamin E analog Size: 50 members Affiliation: Thutewohl, M.; *et al.* [356] Note: Resin attachment through tyrosine phenolic OH at R² or R⁶.

Library: 2.20 Name: Oxime dimer Size: 2565 members Affiliation: Kehoe, J. W.; *et al.* [197] Note: 171-well library with each well containing 2x RCHO and 5x O,Odiaminoalkanediol linkers giving statistical mixture of 15 compounds per well.

Miscellaneous

Library: 2.21 Name: Xanthine derivative Size: *ca*. 18 members Affiliation: Novartis [31]



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k⁵

0 ()₂₋₅ 0 - N

R4

N´]| ö

Library: 2.22 Name: Estradiol derivative Size: 156 members Affiliation: Poirier, D.; *et al.* [298] Note: Sulfamate linker cleaved to give sulfamates or phenols.

R³O

Library: 2.23 Name: 2.5 Diketopiperazinedione Size: 104 members Affiliation: Carbonell, T.; *et al.* [62] Note: Synthesis of initial screening library then a follow-up library.

Library: 2.24 Name: Prolyl/pipecolyl amide Size: 50 members Affiliation: Guilford Pharm. [376]



OH

 $R^3 = H, SO_2NH_2$

NHR¹ \dot{R}^2



Enzyme: Geranylgeranyl transferase(GGTase; C. albicans) Activity: $IC_{50} = 10 \text{ nM}$



Enzyme: Protein farnesyltransferase Activity: IC_{50} = 6.4 μ M



Enzyme: Tyrosylprotein sulfotransferase-2 Activity: IC_{50} = 30 μM



Enzyme: Phosphodiesterase 5 (PDE5) Activity: 90% inhibition at 1 μ M

OH Ĥ 0 H_2N-S

Enzyme: Steroid sulfatase Activity: 98% inhibition at 1 μ M

0 O

Enzyme: Acetylcholinesterase Activity: IC_{50} = 2.2 $\mu M;$ selective versus butyrylcholinesterase

ŚO₂ Ö

Enzyme: FKBP12 Activity: K_i = 640 nM



Table 3. Chemical Libraries Targeting G-Protein Coupled Receptors (Asterisk (*), Point of Attachment to Resin)

In aphabetical order.

Library: 3.1 Name: Cyclopentyladenine Size: *ca*. 20 members Affiliation: Van Calenbergh, S.; *et al.* [364]



Receptor: Adenosine A₁ (human) Activity: K_i = 137 nM



Receptor: α_{1a} Adrenergic Activity: pK_i = 7.36 (non selective agonist)

Library: 3.3 Name: Benzothiadiazide Size: *ca.* 64 members Affiliation: Phillips, D.; *et al.* [295]





Receptor: AMPA (glutamate subgroup) Activity: EC_{50} = 22 μ M



Library: 3.8 Name: Biaryl acylsulfonamide Size: >25 members Affiliation: Merck Frosst [133]

Library: 3.9 Name: Iminodiacetamide Size: *ca*. 600 members Affiliation: Goldberg, J.; *et al.* [139] Note: Solution phase synthesis.





 $\begin{array}{l} \mbox{Receptor: EP}_3 \mbox{ prostanoid receptor} \\ \mbox{Activity: } K_i = 25 \mbox{ nM (antagonist);} \\ \mbox{K}_i = > 1000 \mbox{ nM, EP}_1 \mbox{ and EP}_2 \end{array}$

 $\begin{array}{c} R^{1}HN & \bigcirc & \bigcirc & \bigcirc & \bigcirc & \mathsf{NHR}^{1} \\ & & & & & & \\ R^{2}HN & & & & & & \\ & & & & & & & \\ \end{array}$

ŃН \cap MeO₂C .CO₂Me ŇН ΗŇ 0.0 ö 0 [] 0 ö

Receptor: Erythropoietin (EPOr) Activity: 32% inhibition at 50 μM to displace 125 I-EPO from immobilized EPOr



Library: 3.15 Name: 1-Cyclohexylmethyl-3-aminomethylpiperidine Size: 1000 members Affiliation: Banyu [324]





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solid-phase libraries.

Library: 3.17 Name: Spirohydantoin Size: Not defined Affiliation: Hoffmann-La Roche [42] Note: All library members incorporated 3,5-bis(trifluoromethyl)phenyl, a known NK-1 receptor pharmacophore.

Library: 3.18 Name: Sulfonamide diamine Size: *ca*. 240 members Affiliation: Synaptic Pharm. [129, 180]

Library: 3.19 Name: Oxobenzothiazolin-3-acetic acid amide Size: 300 members Affiliation: Fugisawa [349]

Library: 3.20 Name: 2-Arylindole Size: 128,000 members Affiliation: Merck [380] Note: Mixture library of 320 pools of 400 compounds each.
$$\label{eq:arso_2hn-X-NH-Y-R} \begin{split} & \text{ArSO}_2\text{HN-X-NH-Y-R} \\ & \text{X = cyclic, acyclic spacer;} \\ & \text{Y = CO, CONH, CH}_2 \end{split}$$

CI

ŃH

R

R



Ŕ2

N-R₂

Receptor: Neurokinin-1 (NK-1, human) Activity: pK_i = 7.34



Receptor: Neuropeptide Y5 Activity: $K_i = 12 \text{ nM}$ (antagonist)

 CF_3 C 0

Receptor: Neuropeptide Y5 Activity: $IC_{50} = 0.70$ nM (antagonist)



Receptor: Neurokinin-1 Activity: K_i = 0.8 nM (antagonist)

H ОН N

Receptor: Neuropepticle Y5 Activity: $K_i = 0.8 \text{ nM}$



Receptor: $5-HT_{2a}$ Activity: K_i = 10 nM



Receptor: $5-HT_6$ Activity: K_i = 0.7 nM



Receptor: Mutant vitamin D (VOR(R27-4L)) Activity: EC₅₀ 3.3 nM (agonist)

OH

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Table 4. Chemical Libraries Targeting Non-G-Protein-Coupled Receptors (Asterisk (*), Point of Attachment to Resin)

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Integrins

Library: 4.1 Name: Biphenyl Size: ca. 30 members Affiliation: Bayer AG [362]

Library: 4.2 Name: Peptidomimetic Size: 240 members Affiliation: Abbott [221]





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Receptor: Vitronectin $\alpha_v\beta_3$ Activity: K_i = 0.7 nM\ (antagonist)

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Target: $\alpha_{v}\beta_{3}$ Activity: IC₅₀ = 0.1 nM (antagonist)

Library: 4.3 Name: Biphenylalanine Size: >100 members Affiliation: Genentech [64]





Library: 4.5

Affiliation: Gottschling, D.; et al. [143]

Name: Peptidomimetic

Size: Not defined





Target: $\alpha_4\beta_1$ /VCAM and $\alpha_4\beta_7$ /MAdCAM Activity: $IC_{50} = 10 \text{ nM}, \alpha_4\beta_1$; IC₅₀ = 2.5 nM, $\alpha_4\beta_7$ (antagonist)



Target: $\alpha_4\beta_1$ /VCAM and $\alpha_4\beta_7$ /MAdCAM Activity: IC₅₀ = 12 nM, $\alpha_4\beta_1$; IC₅₀ = 1.1 nM, $\alpha_4\beta_7$ (antagonist)

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Target: Cell adhesion (α4β7/VCAM-1) Activity: Inhibition observed at 12% of medium control.

Library: 4.6 Name: HUN-7293 analog Size: ca. 40 members Affiliation: Chen, Y.; et al. [75]





Target: VCAM-1 Activity: IC₅₀ = 1.6 nM (antagonist); 18x selectivity versus ICAM-1

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Library: 4.9 Name: *N*-Acylphenylalanine Size: 42 members Affiliation: Roche [73]

Library: 4.10 Name: Squaric acid derivative Size: >50 members Affiliation: Celltech [229]

Miscellaneous

Library: 4.11 Name: Tetrapeptide Size: *ca*. 50 members Affiliation: Kipp, R. A.; *et al.* [205] Note: Sequential Aa optimization of AVPI lead.

Library: 4.12 Name: Benzofuran Size: 320 members Affiliation: Bayer [332]



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 $H-Aa_4-Aa_3-Aa_2-Aa_1-XH$ X = O, NH

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OH N H H H H H OH OH OH OMe

Target: VLA-4 ($\alpha_4\beta_1$) Activity: IC₅₀ = 1.2 nM (antagonist)



Target: VLA-4 Activity: IC₅₀ = 200 nM (antagonist)

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Target: B1R3 (bacolovirus IAP repeat) domain of XIAP (X-linked inhibitor of apoptosis protein) Activity: $K_D = 40$ nM



Receptor: Estrogen ER- α Activity: IC₅₀ = 30 nM





ÔAc ŌCOC₆H₅

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Target: A2780 ovarian cancer cells Activity: IC₅₀ = 0.14 µg/mL (10x more active than paclitaxel)

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. ŌCOC₀H₅

OH OAc





Target: Bacterial periplasmic chaperone PapD (*E. coli*) Activity: Pilicide activity demonstrated by surface plasmon resonance



Name: Adenosine derivative Size: *ca.* 35 members Affiliation: Herforth, C.; *et al.* [162] Note: Acylation (R¹CO) of 5-amino nucleoside templates using carboxylic acids activated by Kenner safety-catch linker.



Target: *P. falciparum* Activity: IC_{50} = 1.5 μM N

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triazoles

Table 5. (Continued) Library: 5.22 Name: Aminopiperidine Size: ca. 50 members R¹R²N-L-N Affiliation: Brinner, K. M.; et al. [51] Note: Multiple libraries exploring R1-R3 **P**3 L = linker in combination with discrete analog Microbe: P. falciparum synthesis. Activity: IC₅₀ = 70 nM; IC₅₀ = 7 nM against chloroquinesensitive 3D7 strain. Library: 5.23 Name: Betulinic amide Size: 18 members Affiliation: Pathak, A.; et al. [289] Note: Analogous chemistry carried using ursolic acid as a template to NH_2 generate second 10 member library. $X = OH; HN OH NH_2$ Microbe: P. falciparum Activity: MIC = 10 µg/mL Library: 5.24 R¹ N R² Name: Acyl hydrazide Size: Not defined Affiliation: Caffrey, C. R.; et al. [58] Microbe: Trypanosoma brucei Activity: $IC_{50} = 2 \mu M$ Table 6. Scaffold Derivatization (Asterisk (*), Point of Attachment to Resin) Part A: Solid-Phase NC OBn NHR OR H₂N-OH • Harned, A. M. [159] • Weissberg, A. [378] Celltech [262] • Crich, D. [101] • Lazny, R. [225] • 8 ex; 58-97% • 11 ex; 33-100% • 8 ex; 45-55% • 6 ex; 73-88% • 12 ex; 19-92% • Mitsunobu coupling of ROH to cleavage of resin-bound • from thiopyranoside · conversion of esters to • alkylation of resinhydroxyalkylamides via attached to resin via norbornenyl N-hydroxysuccinimide amines with CNBr bound hydrazones then capture-ROMP-release with transesterification then O to borinate ester linkage H_2NNH_2 N-acyl migration F HC HO -7 ^{*}R¹R²N—Ar нČ • Garcia, J. [135] • Gros, P. [149] • Gravel, M. [145] • BMS [94] • Revell, J. D. [313] • 5 ex; 76-94% • 8 ex; 69-93% >15 ex; good yield • ca. 14 ex; 21-65% • 10 ex; 42-73% Cu(OAc)₂-mediated · from carbamate-· lithiation of resin-bound derivatization of · ionic liquid accelerated linked amino acid 2-pyridylpiperazine then immobilized arylation of aliphatic Suzuki cross coupling of Weinreb amides cleavage with CICO₂Me aryl boronic acids amines and ArB(OH)₃ resin-bound Arl HO HO-P NR¹R² НŠ NH₂ H_2N • Petricci, E. [294] • Katritzky, A. R. [194] • Mourtas, S. [264] • Rinnova, M. [317] Novartis [377] • 14 ex; 53-92% • 8 ex; 65-98% • ca. 40 ex; 30-99% • 3 ex; good yield • 12 ex; 32-96% Oxone-facilitated cleavage · acylation of resin-bound from resin-bound · condensation of resin- Pd-catalyzed amination amines with acylbenzomercapto acids bound amino acid amides of aryl halides on Rink of corresponding resin-bound with R³CHO and (MeO)₂P(O)H thioester

resins





Part B: Solution-Phase (Continued)





Part A: Solid-Phase







Table 8. Monocyclic Synthesis (Asterisk (*), Point of Attachment to Resin)







ureas with Burgess reagent

then addition of R⁴R³NH and intracyclative cleavage

bound 2-aminothiazoles



 assorted heterocyclic furans from resin-bound furfural · C-sulfanylation of resinbound 2-aminothiazoles

resin-bound glycine





Part A: Solid-Phase

NHR² • Wu, C.-Y. [384]

• 12 ex; 80-99% • from soluble polymer-

supported 4-fluoro-3nitrobenzoic acid



• SIDDCO [280] • 7 ex; 11-20%

• from resin-bound amines and 2-nitrobenzoic acid: final cyclization performed after cleavage



• Macleod, C. [247] • 12 ex; 58-72%

• conversion of resin-bound esters to enol ethers via Ti(IV) benzylidenes bearing a masked nitrogen nucleophile



• BMS [370]

• 15 ex; 0-88% • reductive amination of anilines onto Bal resin

cyclization and cleavage

acylation, then intramolecular



• Hwang, S. H. [178]

• 8 ex; 6-24% yield • use of traceless sulfone linker



- ArQule [155]
- 5 ex; good yield
- from resin-bound chalcone N-methylisatin and fluorophenyl

glycine



- Holland, R. J. [168] • 11 ex; >80%
- · reductive cyclization of
- 2,4,6-substituted-3,5-difluoro nitrobenzenes



• Akamatsu, H. [7]

• 8 ex; 90%

· resin-bound bromomethyl amides alkylated with heterocyclic diamines the condensation with ArCHO



• Liao, Y. [230]

- 90 members
- carbonylative annulation
- of o-alkynylphenols

Part A: Solid-Phase (Continued)



• from resin-bound o-nitrobenzoic acid

· acylation of resin-bound diamine with 4-fluoro-3-nitrobenzioc acid, then multi-step elaboration to dihydroimidazoles

• chiral nitrone addition to resin-bound acrylate; solutionphase example also given

 ca. 23 members • Ugi 4CC







• ca. 7 ex; 34-48%

- DDQ-promoted oxidation
- of phenolic Schiff bases

• 11 ex; 74-96%

• 80 members

strategy

Ugi/de-BOC/cyclize

Cu-catalyzed 3CC



- Barthelemy, S. [21]
- 10 ex; 56-94%

• cyclization o-azido arylimides with perfluoro tagged Ph₃P or resin-bound triarylphosphine



- Organ, M. G. [285]
- 2 ex
- multi-step sequence using RCM chemistry

Part B: Solution-Phase (Continued)



Table 10. Polycyclic and Macrocyclic Synthesis (Asterisk (*), Point of Attachment to Resin)



recyclable acylating

reagent







recyclable RCM catalyst

· universal isonitrile for

Ugi reactions



· acid chloride synthesis

amine acylation



- immobilization of boronic acids
- immobilization of ketones immobilization of olefins for alkylation





 polymer-supported SES group for amine protection





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