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Reviews

Comprehensive Survey of Chemical Libraries for Drug Discovery and Chemical Biology: 2008

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This is the twelfth installment of the comprehensive survey series in combinatorial chemistry.¹ Biologically active libraries reported in 2008 are captured in Tables 1-5 under the headings of proteases, nonproteolytic enzymes, GPCRs, nonGPCRs, and oncolytics/antiinfectives. Table 6 lists molecular probes. Compound collections without disclosed biological activity are delineated in Tables 7-10 under the headings of scaffold derivatization/acyclic synthesis, monocyclic-, bicyclic/spirocyclic-, and polycyclic/macrocyclic synthesis. Polymer-supported reagents/scavengers/linkers are presented in Tables 11 (nonfluorous) and Table 12 (fluorous). There are 505 libraries and 30 molecular probes extracted from 490 literature citations.^{2–491} Approximately 90% of the citations originated from academic laboratories and $\sim 80\%$ of the chemical libraries relied on solution-phase synthesis for their preparation.

There are 26 vignettes in the 2008 literature review. Selected vignettes on biologically active libraries include HIV-1 nonnucleoside reverse transcriptase inhibitors,²⁷⁷ Aurora A kinase inhibitors,⁷² orexin-2 receptor antagonists,¹⁰⁴ mGluR4 allosteric modulators,²⁸⁵ p38a kinase inhibitors,³⁵ TACE inhibitors,²⁸³ cannabinoid-2 receptor agonists,⁴¹² CCR2b receptor antagonists,²⁷⁵ and DOSderived MRSA growth inhibitors.³³⁷ The molecular probe selections include: dapagliflozin,⁴⁸⁶ dorsomorphin,⁴⁷³ a neuregulin/ErbB4 signaling inhibitor,474 and proteinase activated receptor-2 (PAR-2) agonist AC-262613.468 Highthroughput chemical methodology selections include: traceless gem-difluorinating cleavage of resin-bound dithianes,⁴⁰⁸ traceless solid-phase synthesis of thiomorpholinones³³⁴ and 1,4-diazepinones,³³³ N-hydroxyethyl-1,4-diazepinone synthesis, and parallel derivatization,¹⁵⁰ multicomponent reactions (MCR) yielding spiroquinolines,⁴⁰ tricyclics,¹¹⁷ 1,2-amino alcohols,¹⁰¹ and α -amino acid amides (unprotected N-terminus) by a novel phenylphosphinic acid-catalyzed Ugi reaction,²⁹⁶ parallel synthesis of 1,2,4-oxadiazoles via a continuous microfluidic flow reaction sequence,¹⁶⁷ and the use of fluorous technology in natural product mixture synthesis²⁰⁵ and in the synthesis of β , β -difluoro- α -amino acid derivatives,⁴⁵⁸ natural product-like diols,459 and arrays of substituted 4-thiazolidinones and 4-thiazinanones.440

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Related publications and reviews appeared in 2008 on microwave-assisted C–C and C–N bond formation, ⁴⁹² multicomponent, cascade, and tandem reactions, ^{493–497} high-throughput heterocyclic synthesis, ^{498–505} click chemistry, ^{506–508} dynamic combinatorial chemistry, ^{509–511} diversity-oriented synthesis (DOS), ^{512–514} library automation and analysis, ^{515–524} library design, ^{525–530} diversity analysis and evaluation, ^{531–537} chemical genomics, ^{538–544} colorimetric tests for the detection of amines and thiols, ⁵⁴⁵ alkylsquarate scaffold synthesis, ⁵⁴⁶ color-based encoding, ⁵⁴⁷ new photolabile tags, ⁵⁴⁸ continuous-flow reactors and application in synthesis, ^{549–551} strategies for early drug discovery, ^{552–554} combinatorial libraries against cancer and protease targets, ⁵⁵⁵ strategies for mixture-based synthetic combinatorial libraries, ⁵⁵⁶ libraries from libraries, ⁵⁵⁷ water-accelerated synthetic reactions, ⁵⁵⁸ estimation of protein–ligand affinity by NMR, ⁵⁵⁹ metal-catalyzed C–C bond formation in SPOS, ⁵⁶⁰ affinity selection-mass spectrometry in HTS, ⁵⁶¹ and fluorous chemistry and separa-

tions. 562-566

HIV-1 Wild-Type and Mutant Nonnucleoside Reverse Transcriptase Inhibitors.²⁷⁷ Human immunodeficiency virus (HIV) is a retrovirus causing acquired immunodeficiency syndrome (AIDS). There are two types of HIV, HIV-1 and HIV-2, where the former is highly virulent. Current therapies to control HIV-1 include 22 FDAapproved agents targeting virus-cell fusion, viral protease, and viral reverse transcriptase (RT). The latter target class is divided into two categories: (a) nucleoside and nucleotide analog RT inhibitors (NRTIs) and (b) nonnucleoside RT inhibitors (NNRTIs). Compared to NRTIs, NNRTIs are more specific and exhibit less severe adverse effects; however, these inhibitors are rendered ineffective upon relatively minor HIV-1 RT mutations.

Botta and colleagues previously identified dihydroalkoxybenzyloxopyrimidines (DABOs) as potent NNRTIs.⁵⁶⁷ Although their initial research led to the discovery of compounds exhibiting nano- and subnanomolar anti-HIV potency in lymphoid cells, these compounds were not as effective against mutant strains. To identify novel anti-HIV wild-type and mutant RT inhibitors, Mugnaini and colleagues designed a series of arylalkylthio substituted DABOs 5 (Figure 1).²⁷⁷ The parallel solution-phase synthetic approach began with potassium ethyl 2-substituted malonates 1 which are reacted with MgCl₂ and several phenyl acetic acids 2 to afford β -ketoesters 3. Treatment of **3** with thiourea in the presence of sodium ethoxide produced uracils 4 which in turn were thio alkylated under microwave irradiation conditions with either benzyl/alkyl halides or alcohols under Mitsunobu conditions to afford the desired sulfur-substituted heterocycles 5 (Library 2.37).

Initial structure—activity relationships (SAR) showed a very distinct pattern in which a benzyl group at position 6 was essential for activity and that *para*-substitution on this group was detrimental to activity. With this nascent SAR, the authors proceeded to further explore substitution effects on the benzyl group, determining that 2,6-dichloro and 2,6-difluoro substitutions were favorable for activity. Thiobenzylated analogs (6-11, 14-19) exhibited submicromolar potency in both enzyme and cell-based (NL4-3 cells) assays. Substitution studies on the aromatic unit of the S-benzyl moiety (R⁴) indicated that para-substitution was highly beneficial to activity regardless of the electronic nature of the substituent (e.g., 6-8, 4-OMe, 4-CN, 4-NO₂; ID₅₀ = 0.032, 0.1, 0.075 μ M, respectively). A 2-carbon chain between the phenyl and thio groups was tolerated, but a 3-carbon chain was detrimental to activity (12 versus 13, $ID_{50} = 0.07$ and 9.76 μ M, respectively). Small substituents on the oxopyrimidine moiety were also well tolerated (R¹: 6, 9, ethyl and methyl; $ID_{50} = 0.032$ and 0.77 μ M, respectively). Lastly, it was determined that substitution on the benzyl carbon (\mathbb{R}^2) was well tolerated with a trend of exhibiting higher activity when substituents were present (14, 17, $ID_{50} = 0.03$ and $0.003 \ \mu M$, respectively). From this S-benzyl analog class, compound 17 was the most potent inhibitor: $ID_{50} = 3 \text{ nM}$; $EC_{50} =$ 25 pM in NL4-3 wild-type cells. This agent possessed submicromolar activity against three mutant strains (K103N, Y181C, Y188L; $EC_{50} = 0.55$, 0.3, and 0.95 μ M, respectively).

While changing the benzyl methylene unit for a vinyl unit was not favored (**20**, $ID_{50} = 1.98 \,\mu$ M), an allyl group led to increased activity (**21**, $ID_{50} = 0.007 \,\mu$ M). Among the 3-arylallylphenyl analogs (**21–28**) sharing many of the substituents present in **17**, analog **27** was found to exhibit optimal anti-HIV activity in both the enzymatic and cellbased assays ($ID_{50} = 9 \,n$ M; $EC_{50} = 0.4 \,n$ M). Interestingly, exchanging the allyl group in **27** for a propargyl group was detrimental to activity, particularly in NL4–3 wild-type cells.

Although compounds **29–31** having a *para*-methoxybenzyloxy ethyl group linked to the sulfur atom were potent enzyme inhibitors, it was only 30 that exhibited activity against the wild-type and the K103N, Y181C, and Y188L mutant strains (EC₅₀ = 0.0011, 0.92, 0.65, and $0.32 \,\mu\text{M}$, respectively). To rationalize these observations, the authors used molecular docking in conjunction with crystallographic data for reverse transcriptase (RT)-ligands of the K103N, Y181C, Y188L mutant strains. Docking studies with the Y188L mutant correctly predicted inhibitors 17-19 to be the most potent. The *R*-enantiomer of 17 adopted a similar orientation to 32, where the pyrimidinone ring of 17 occupied the same region as 32's quinoxaline ring. Inhibitor 17 also formed two hydrogen bonding interactions with the CO and NH groups of Lys101 in the backbone, the same hydrogen bonds found in 32. Moreover, the para-methoxybenzyl group was positioned in a large hydrophobic-reach pocket defined by Val106, Pro225, Pro236, and Phe227, justifying the role and interactions of the para-methoxybenzyl group with the protein. These docking parameters, however, could not fully explain the potent activity exhibited by compound 30.

Tethering Dynamic Combinatorial Chemistry: Aurora A Kinase Inhibitors.⁷² Aurora kinases are serine/threonine kinases involved in mitosis, particularly in chromosome segregation and cell division, and their functional deregu-



Figure 1. HIV-1 nonnucleoside reverse transcriptase inhibitors.²⁷⁷

Me

Me

Me

2.6-diF

2,6-diF

2,6-diF

4-NO2

4-OMe

4-CN

0.016

0.009

0.048

0.0002

0.0004

0.59

26

27

28

1

1

1

Et

Et

Et

lation results in tumorigenesis. There are three known mammalian Aurora isozymes A, B, and C. Researchers at Sunesis Pharmaceuticals used site-specific dynamic combinatorial chemistry as an approach to the discovery of Aurora A kinase inhibitors. A unique tethering technology was developed in which a cysteine residue near a site of interest on a protein target is used to covalently bind with small thiol-containing molecular fragments via a disulfide bond in order to identify weak binding fragments. The second generation tethering technology incorporated the use of "extenders" or bifunctional molecular fragments containing an electrophilic unit and a thioacetate unit that are covalently linked to a protein's cysteine residue and can also interact reversibly with other fragments that bind at a neighboring site. A new third-generation tethering approach employs "dynamic extenders", which contain two disulfide units: one covalently binds with a cysteine

Antiviral activity and SAR (1)







residue (disulfide linkage), while the second disulfide unit reversibly captures other fragments. This third-generation tethering approach affords a combination of dynamic combinatorial chemistry with fragment-based lead discovery.

This new technology was used to identify novel binding fragments at the ATP catalytic site of Aurora A (Figure 2). The researchers first introduced a cysteine residue (T217C) near the ATP pocket via site-directed mutagenesis. A collection of approximately 4500 disulfidecontaining fragments was screened in pools of 10 against the mutated Aurora A protein under partially reducing conditions in the presence of the diaminopyrimidine-based dynamic extender **33** (Library 2.2). Hits formed a stable complex with the T217C-mutated Aurora A kinase and were identified using electrospray ionization mass spectrometry. The extender-fragment combinations were elucidated and converted into the respective nondisulfide









Figure 2. Sunesis' tethering dynamic combinatorial approach to Aurora A kinase inhibitors.⁷²

small molecules, affording **34** (IC₅₀ = 17 μ M). Further experiments identified a purine-based dynamic extender affording **35** (IC₅₀ = 2.9 μ M), an analog 5-fold more potent than **34**. The results were validated by cocrystallizing **35** with nonmutated (wild-type) Aurora A kinase. As observed in the X-ray cocrystal structure, the purine unit of **35** is accommodated at the ATP binding pocket, the phenyl rings form a π -cationic interaction with the side chain nitrogen of residue Lys162 and the distal amide



43: OX2-R pIC₅₀ = 7.29

Figure 3. Orexin-2 receptor antagonists.¹⁰⁴

group interacts with the backbone of residue Phe144 via a hydrogen-bond interaction.

OX2 Receptor Antagonists.¹⁰⁴ The orexins (orexin A and orexin B, also named hypocretin 1 and 2, respectively) are hypothalamic peptides playing an important role in the regulation of the sleep-wake cycle, feeding behavior, regulation of gastric acid secretion, metabolic rate and





- Potentiate the EC20 of glutamate in mGluR4/Gqi5 CHO cells: EC_{50} = 4.6 μ M - No effect on mGluR4 in the absence of

- glutamate
- Selective versus mGluR5
- Full mGluR1 antagonist (IC₅₀ = 2.6 μM)
- Poor metabolic stability

Library synthesis and inputs:



Figure 4. mGluR4 positive allosteric modulators.²⁸⁵

blood pressure. The orexins are endogeneous ligands of two receptors, OX1-R and OX2-R, which belong to the superfamily of G-protein coupled receptors (GPCRs). While OX2-R binds both neuropeptides with similar affinity, OX1-R binds orexin A with slightly higher affinity than orexin B. Recent data suggests that OX2-R is associated with the sleep/wake cycle and that mutations of this receptor are related to canine narcolepsy. Human narcolepsy also appears to be linked to deficient orexin signaling, most likely related to immune ablation of

Kinase ATP-domain binding model and library 2.12 design



Library synthesis and representative actives



Figure 5. p38α kinase inhibitors.³⁵

orexinergic neurons in the lateral hypothalamus. With these findings, the search for selective OX2-R antagonists for the treatment of sleep and neurological disorders has recently increased in intensity in the pharmaceutical industry. Researchers at Pharmacopeia conducted a high throughput screen of 90 distinct ECLiPS (Encoded Combinatorial Libraries on Polymeric Support) libraries to identify novel OX2-R antagonists (Figure 3). Libraries containing over four million compounds were screened at the human OX2-R using a Fluorescent Imaging Plate Reader (FLIPR)-based calcium assay. Active hits were found from a subset library (~140 000 members) of general structure 36. In particular, (3,4-dimethoxyphenoxy)-alkylaminoacetamides 41 were identified from this screening campaign as a novel chemical class of potent OX2-R antagonists. The combinatorial library synthesis of 41 (Library 3.27) is outlined in Figure 3. Condensation of primary amines (R^1NH_2) with resin 37, prepared from Tentagel polymer support and a photolabile linker, provided the corresponding amine-bound resin 38. This resin was converted to amino alcohol resin 39 by condensation with bromoacetic acids followed by addition of amines (R^2NH_2). Alkylation of the amine functionality of **39** under reductive amination conditions, followed by Mitsunobu coupling of the terminal alcohol with various phenolic derivatives, gave resin 40 which was cleaved by irradiation to release the desired library compounds 41. SAR analysis in this series demonstrated that isobutyl, benzyl, and 2-chlorobenzyl groups were preferred at the R^1 position, while the R^2 position could accommodate a wide variety of substituents. For the more potent analogs, the α -aminoamide moiety of **41** was linked to the key 3,4-dimethoxyphenoxy pharmacophoric group by ethylene, α -methylethylene, β -methylethylene, or propylene linkers. From this work, compounds 42 and 43 were identified as potent OX2-R antagonists (pIC₅₀ = 7.71 and 7.29, respectively).

mGluR4 Positive Allosteric Modulators.²⁸⁵ The majority of drugs that target GPCRs interact with the receptor orthosteric site (i.e., the same domain recognized by the endogenous agonist for the receptor). Recently, several research groups identified ligands binding to allosteric sites of GPCRs (i.e., binding domains topographically distinct from the orthosteric sites). The binding of a ligand to an allosteric site causes a conformational change in the receptor protein that is transmitted to the orthosteric site (and vice versa). Allosteric ligands are also called allosteric modulators because they can alter the binding affinity or the signal produced to the cell by an orthosteric ligand. One of the main research interests in the Conn group at Vanderbilt University is to identify allosteric modulators of various proteins, including the metabotropic glutamate receptor subtype 4 (mGluR4). L-Glutamate, the major excitatory amino acid neurotransmitter in the CNS, binds to and activates several classes of receptors divided into two groups, the ionotropic (iGluR) and metabotropic glutamate receptors (mGluR). The metabotropic glutamate receptors, belonging to the family of GPCRs, have been divided into eight subtypes based on sequence homology (group I mGluR1, mGluR5; group II mGluR2, mGluR3; group III mGluR4, mGluR6, mGluR7, mGluR8). Preclinical data support the utility of mGluR4 agonists in various indications including anxiety, depression, and Parkinson disease. A high throughput screening campaign of Vanderbilt's compound collection against mGluR4 identified the pyrazolo[3,4-d]pyrimidine 44 as a positive allosteric modulator (Figure 4). Compound 44 produced an EC₅₀ for potentiation of the agonist (glutamate) response of 4.6 μ M. Importantly, this compound did not produce any effect on mGluR4 activity in the absence of glutamate.

Compound 44 also produced a much greater shift of the glutamate concentration-response curve when compared to a literature referenced mGluR4 positive allosteric modulator. To support the hit-to-lead campaign, a 126member library of substituted pyrazolo[3,4-d]pyrimidines 45 (Library 3.9) was prepared. A general microwaveassisted protocol to this heterocyclic system was developed to support parallel synthesis. Hence, treatment of aryl hydrazines with ethoxymethylenemalonitrile 46 in ethanol (MW, 105 °C, 10 min) yielded 4-cyano-5-aminopyrazole 47 in 54-77% yields. Hydrolysis of the nitrile with aqueous H₂SO₄ gave the corresponding carboxamides 48 which were converted to the 1-aryl-*1H*-pyrazolo[3,4d]pyrimidin-4-ols 49 in neat formamide (MW, 200 °C, 20 min). Microwave-assisted-conversion of 49 to the corresponding heteroaryl chloride with POCl₃ and subsequent S_NAr with various amines provided pyrazolo[3,4d]pyrimidines 45. A majority of the analogs prepared did not show any activity as mGluR4 positive allosteric modulators. The "flat" SAR observed in this series has also been an issue for other chemical series of mGluR positive allosteric modulators.

Kinase Inhibitors.³⁵ Protein kinases constitute one of the largest protein families in humans. These kinases regulate the majority of signal transduction pathways in cells, and thus play an important role in cell growth, metabolism, differentiation, and apoptosis. Deregulation of protein kinases is implicated in a number of diseases, including cancer, diabetes, and inflammation. For these reasons, targeted inhibition of protein kinases is an attractive therapeutic strategy. All kinase enzymes share a catalytic domain that contains a cleft where adenosine triphosphate (ATP) binds. This catalytic cleft is a major focus of small molecule drug design. Functional states of a typical protein kinase can be characterized by the position of a conserved DFG (aspartate, phenylalanine, glycine) motif in its activation loop. The majority of inhibitors target the ATP site of the kinase in its active "DFG-in" state. In contrast, the so-called type-II inhibitors induce a distinct "DFG-out" conformation and occupy an additional hydrophobic pocket created by this rearrangement. These type-II inhibitors possess several advantages over ATP-site compounds, including improved kinase selectivity and slower off-rates. Researchers at Glaxo-SmithKline recently identified a series of biphenyl amides as p38a mitogen-activated protein (MAP) kinase inhibitors (Figure 5).³⁵ Representatives of these agents are compounds 50 and 51, binding to the p38 α MAP kinase in either the DFG-in (compound 50) or DFG-out (compound 51) conformation. To further explore the SAR in this series, a 172-member biaryl library 52 (Library 2.12) was prepared in one step (Suzuki coupling) and profiled in the p38 α competitive binding assay. The R¹ substituents (A-D) targeting the lipophilic interior of the ATP-site (or back-pocket) were selected based on the structure of other potent kinase inhibitors. The R² substituents, designed to occupy the purine subpocket, were chosen based on information obtained from the crystal structures of 50 and **51** binding to p38α MAP kinase. A total of 87 compounds,



Figure 6. TACE inhibitors.²⁸³

including 60 different purine subpocket-binding groups, displayed K_i affinity at p38 α of less than 100 nM. Selected compounds were also profiled against other kinases including Raf/Mek/Erk, CDK2, cFMS, EGFR, ErbB4, JNK3, LCK, PLK1, SGK1 and VEGFR2. This study led to the identification of compound **56**, a potent p38 α MAP kinase inhibitor displaying >20-fold selectivity over the other kinases profiled. Additional studies showed that **56**

inhibited TNF α production in peripheral blood mononuclear cells (IC₅₀ = 100 nM) and human whole blood (IC₅₀ = 1000 nM). This compound also demonstrated favorable pharmacokinetic properties in rat. In contrast to compounds from arrays **A**-**C**, compounds from array **D**, such as **57** displayed potent affinity for cFMS and LCK, but weak affinity for p38 α .

TACE Inhibitors.²⁸³ The overexpression of the proinflammatory cytokine tumor necrosis factor- α (TNF- α) has been implicated in numerous pathological conditions. The anti-TNF- α biological therapeutics entaneracept, infliximab, and adalimumab have demonstrated clinical success in treating inflammatory and autoimmune diseases validating the modulation of TNF- α as a drug discovery target. Inhibition of TNF- α converting enzyme (TACE) is considered an attractive mechanism to control the release of TNF- α and a viable therapy for the treatment of rheumatoid arthritis and Crohn's disease, as these conditions are caused by overexpression of TNF-a. TACE (also called ADAM-17) is a Zn-dependent metalloproteinase belonging to a subclass of the metzincin family. Because of structure similarities between the active sites of TACE and related matrix metalloproteinases (MMPs), early TACE inhibitors were derived from MMP inhibitors and suffered from broad spectrum activity against the MMP family. Nonselective TACE inhibitors have been shown to produce musculoskeletal side effects in clinical trials. Research efforts are currently concentrated on identifying TACE inhibitors with high selectivity over the other MMPs. Researchers at Schering-Plough designed a novel series of TACE inhibitors based on the information obtained from the X-ray crystal structure of the known TACE inhibitor 58 docked into the enzyme active site. In the newly designed structures 59 (Figure 6), the key hydroxamate moiety can coordinate the zinc atom of the enzyme in a bidentate fashion, while the carbonyl moiety can potentially participate in hydrogen-bonding interactions with the Leu348 and Gly349 amino acid residues of the enzyme. An additional lipophilic moiety is necessary to occupy the lipophilic S1'-S3' binding pocket. Four compounds were prepared as proof of concept molecules for this study. Two of these compounds, 60 and 61, were found to bind to TACE with submicromolar affinity (60: $K_i = 500 \text{ nM}; 61: K_i = 800 \text{ nM}).$ Based on these encouraging results, analogs of 60 and 61 were prepared using solution-phase and solid-phase synthesis (Library 1.4). The preparation of selected examples is shown in Figure 6. Cyclopropanation of the acrylate derivative 62 in the presence of *tert*-butyloxymethyltetrahydrothiophenium bromide provided the cyclopropane derivative 63, which was converted to the chiral derivative 64 by hydrogenation, followed by chiral separation of the corresponding racemic mixture. Coupling of the acyl chloride 65 (obtained in three steps from 64) to the resinbound hydroxylamine 67, followed by cleavage of the acetyl protecting group provided the resin-bound phenol **68**. Alkylation or arylation of the phenolic hydroxyl group of 68 followed by TFA cleavage and HPLC purification provided the desired compounds 69 and 70. Loading of 71 (prepared in 2 steps from 64) to the hydroxylamine resin provided the corresponding hydroxamate resin, which was converted to 72 by subsequent ester hydrolysis. Coupling of the carboxylic acid moiety of 72 with either isopropyl alcohol or benzyl amine followed by TFA cleavage afforded the corresponding isopropyl ester 73 and benzylamide 74, respectively. Multiple derivatives from both series were identified as potent TACE inhibitors. In particular, compounds 75 and 76 displayed K_i values for TACE of 12 nM and 8 nM, respectively. Despite the close structural similarity between these two compounds, their selectivity profile for TACE versus other members of the MMP family was surprisingly quite different. Hence, while 75 displayed poor selectivity over ADAM-10, compound 76 was identified as a selective TACE inhibitor. X-ray crystallography of 75 and 76 binding to TACE revealed two different binding modes for these inhibitors. This could account for the difference in SAR trends and selectivity profiles among the two series of inhibitors.

Cannabinoid 2 Receptor Agonists.⁴¹² There are three cannabinoid receptors linked to nociception and agonists of the receptors are thought to be potential analgesics. Worm and colleagues describe the synthesis of a series of sulfamoyl benzamides for use as CB₂ agonists. On the basis of the high-throughput screening hit 77, the authors investigated three points of derivatization for the sulfobenzoic acid core (Figure 7) using a combination of parallel and solid-phase synthetic approaches. Coupling of amines **81a**-**h** with the commercially available sulforyl chloride **79** gave the sulfonamides **80**. A diverse set of resin bound amines was prepared by reductive amination of amines to an aldehyde-based polystyrene resin. These were subsequently coupled to the sulfonamides. Cleavage of the resin with TFA gave the desired compounds 78 (Library 3.4). Approximately half of the compounds produced in this library led to retention or increased affinity for the CB₂ receptor when compared to 77, while the remaining compounds exhibited reduced affinity for both CB receptors. In an effort to improve affinity for the CB₂ receptor, the authors then looked to introduce highly branched amines (83a-h) at the R³ position. Initial attempts to prepare the amides using the resin bound strategy previously employed failed and these amides were subsequently prepared via solution phase methodology. Introduction of the S-fenchyl group (84) led to vastly improved affinity for the CB₂ receptor. Replacement of piperidine with morpholine at the R^2 position (85) led to a further increase in CB₂ receptor affinity. Substitution of the bromo substituent at R^1 with a methyl group (86) had no significant effect on binding affinity for the CB₂ receptor. However, inversion of the fenchylamine stereocenter at R^3 (87) led to a dramatic decrease in CB_2 receptor affinity. All compounds were then tested in a $[^{35}S]GTP\gamma S$ functional assay. All were found to be full agonists, with the exception of 90, which exhibited inverse agonist activity.

CCR2b Antagonists.²⁷⁵ In an extension of their previous research, Moree and co-workers report a new set of potent



Figure 7. Adolor's CB2 agonists.⁴¹²

CCR2b antagonists.²⁷⁵ Based on lead compound **91**, the authors identified three areas of derivatization for SAR development (Figure 8). Initial research to investigate the R^{1} functionality was performed in solution phase. Representative compounds include 95-100. A clear preference for *para*-substituted benzyl substituents was established through comparison of the CCR2b affinity of compound 91 with compounds 96 and 97. Expansion of the range of 4-benzyl substituents showed a preference for small electron donating groups (98) over electron withdrawing moieties (99). Interestingly, 2,3- or 2,4- disubstituted benzyl groups were also tolerated and in some instances led to enhanced affinity for the CCR2b receptor (100). Simultaneous modification of the R² and R³ substituents was explored while holding the R^1 substituent (4-chlorobenzyl) constant. Representative compounds include

103–108. Utilizing a variety of α and β amino acids, it was established that even small modifications to the amino acid residue (103, 104) led to significantly reduced affinity for the receptor. Likewise, relocation of the R^2 substituent from the meta to the ortho (105) or para (106) positions led to decreased receptor affinity. However, 2,5-disubstitution of the benzyl ring led to compounds with much greater affinity for the CCR2b receptor. Following this finding, attention was refocused on the R^1 substituent to further optimize the ligands. Keeping the 2-amino-5trifluoromethylbenzamide R² substituent constant, it was found that modification of the R¹ substituent through the use of mono substituted benzyl compounds offered very little improvement in receptor binding. However, disubstituted analogs (109, 110) led to compounds with single digit nanomolar affinity for the CCR2b receptor and potent



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MCP-1 induced chemotaxic activity. During the execution of the research, a solid-phase strategy for simultaneous derivatization of the R¹ and R² substituents was devised. Using the acid sensitive methoxybenzaldehyde (AMEBA) resin, the protected aminopyrrolidine was attached to the preloaded amino acid of interest. Introduction of the R² substituents was then achieved through amide formation with a benzoic acid. Removal of the Fmoc protection and reductive amination with a variety of benzaldehydes afforded the resin bound products. Cleavage of the resin using TFA gave the target compounds, which were then purified using ion exchange chromatography.

Anti-MRSA Agent Discovery Using Diversity-Oriented Synthesis (DOS).³⁷⁷ Bacterial drug resistance is the cause of serious health problems. Of particular concern is the rise of nocosomial infections in patients exposed to drug-resistant bacteria in hospitals and day-care centers. Drug-resistant bacteria, also referred to as "superbugs", cannot be treated successfully with available antibacterials and may lead to mortality within a week of infection. One of the most devastating superbugs is methicillin-resistance *Staphylococcus aureus* (MRSA). MRSA is resistant to \sim 30 different antibiotics.

Thomas and colleagues employed DOS to generate Library 5.30 derived from 18 different natural-productlike scaffolds (115-132) to discover novel compounds with anti-MRSA activity (Figure 9). A silyl-based solidphase synthesis strategy based on resin 113 was pursued for three reasons: (1) solid-phase simplifies optimization and workup procedures, (2) the phosphonate moiety in 113 allows for the stereoselective formation of α,β unsaturated acyl imidazolidinones to introduce selected stereofunctionalities, and (3) the imidazolidinone linker can be hydrolyzed, esterified, reduced, and amidated to produce desired molecular diversity. Resin 113 was exposed to alkyl, aryl, and heteroaryl aldehydes in the presence of LiBr and DBU to afford a set of twelve α,β unsaturated acyl imidazolidinones 114. Building blocks 114 were subjected to three enantioselective skeletal diversification reactions creating three primary DOS spurs: [2 + 3] cycloaddition (114 \rightarrow pyrrolidines 115); dihydroxylation (114 \rightarrow diols 118); and the Diels-Alder cycloaddition (114 \rightarrow norbornenes 119). Each spur was then submitted to further derivatization. Acylation and reductive amination of pyrrolidine spur 115 afforded 116 and 117, respectively. Spur 118 was sequentially treated with (a) $SOCl_2$ (dioxathiolane formation), (b) NaN_3 (azido alcohol intermediate), and (c) dimethyl acetyldicarboxylate ([3+2] cyclization) yielding triazoles **120**. Alternatively, 118 was O-alkylated, condensed with ketones and aldehydes generating diether, ketal, and hemiacetal analogs $(119 \rightarrow 121, 122, 123)$. Utilizing oxidation and metathesis reactions, norbornene spur 119 was diversified to bicyclic and tricyclic scaffolds 124-128. A secondary spur 126 was obtained through robust synthetic transformations furnishing five additional scaffolds 129-132. After resin cleavage, 242 compounds were obtained in 1-20 mgquantities (purified as necessary to ensure >90% purity), and all characterized by ¹H NMR and LCMS. To

determine its overall diversity, library 5.30 was evaluated against a series of "benchmark" collections including biologically active molecules in the MDL Drug Data Report database (MDDR). Diversity was assessed through principal component analysis (PCA). Library 5.30 was screened against three S. aureus strains: methicillinsusceptible S. aureus (MSSA), and the penicillin- and erythromycin-resistant U.K. epidemic methicillin-resistant strains EMRSA 15 and EMRSA 16. Compounds 133 and 134 were identified as hits derived from spur 119 \rightarrow bicyclic amine family 129. The nascent SAR (13 analogs) around the more potent compound 133 named (-)gemmacin indicated that both carboxylic and nitro groups are essential for activity. (-)-Gemmacin exhibited higher selectivity toward Gram-positive over Gram-negative bacteria, low antifungal activity (MIC₅₀ > 64 mg/mL in 7 *Candida* species) and low mammalian cell toxicity: MIC₅₀ > 64 mg/mL in human epithelial cells.

Molecular Probe: SGLT2 Inhibitor Dapagliflozin.⁴⁸⁶ A number of important molecules have recently joined in the fight against type 2 diabetes. Included among their ranks is dapagliflozin 137 (Figure 10), a recently reported inhibitor of renal sodium-dependent glucose cotransporter 2 (SGLT2).⁴⁸⁶ The SGLTs regulate the reabsorption of plasma glucose and SGLT2 is responsible for this process in the kidney. Inhibition of SGLT2 is theorized to be a unique method for control of plasma glucose levels by promoting glucose excretion. Selective inhibitors of SGLT2 may minimize side effects associated with SGLT1 inhibition in the gastrointestinal tract including diarrhea and dehydration. Furthermore, controlling glucose levels via this mechanism would not alter the native regulatory machinery for glucose, thus avoiding issues of hypoglycemia. The natural product O-glucoside phlorizin 135 has been demonstrated to inhibit SGLTs in a nonspecific manner and is a well-known glucosuric agent. Given its lack of selectivity and poor bioavailability, researchers at Bristol-Myers Squibb Company opted to use the core structure of phlorizin as a starting point for the rational design of a novel, selective inhibitor of SGLT2. A key component of their efforts was the switch from Oglycosides to more robust C-glycosides. From these efforts they found the C-aryl glycoside **136** with modest activity versus SGLT2. Exploration of this scaffold eventually led to dapagliflozin 137. Synthetically, the core C-aryl glycoside is established via lithium-bromine exchange on the appropriately substituted aromatic precursor and nucleophilic attack on a persilylated gluconolactone. This reaction gives a mixture of lactols that requires in situ conversion to the desilylated O-methyl lactol. Reduction of the anomeric methoxy was accomplished with triethylsilane and $BF_3 \cdot OEt_2$ and peracetylation allows a key crystallization which is necessary to ensure collection of the β -anomer. A final hydrolysis provided dapagliflozin. Biochemically, dapagliflozin inhibits SGLT2 with an EC₅₀ of 1.1 nM and SGLT1 with an EC_{50} of 1.39 μM (1200 fold selectivity for SGLT2). In vivo analysis in rats (both glucosuric response and mean blood glucose levels) was promising and dapagliflozin was advanced for clinical



Figure 9. DOS library of MRSA growth inhibitors.377

evaluation. Dapagliflozin 137, a probe with a unique chemotype that cleverly incorporates a carbohydrate scaffold, represents an important new tool in the fight against type 2 diabetes.

Molecular Probe: BMP Signaling Inhibitor Dorsomorphin.⁴⁷³ The use of small molecules to manipulate stem cells (adult and embryonic), reprogram lineage committed cells or revert defined cell types to a pluripotent state has



Figure 10. Discovery of dapagliflozin.486

AcC

been the focus of much effort in recent years.⁵⁶⁸ Many of these studies are forward chemical genomics studies that define molecules with desired phenotypes without understanding the mechanism behind these actions. However, advances in the study of the endogenous signals that control and regulate the fate of multipotent cells during development has provided a plethora of new information regarding important targets that control cellular differentiation. These studies allow reverse chemical genomics screens aimed at defining potent and selective molecules that modulate important cellular targets and provide researchers novel tools for this rapidly advancing field. An impressive example of this is the discovery of dorsomorphin 138 (Figure 11), a small molecule inhibitor of bone morphogenetic protein (BMP) signaling. BMP signaling controls a number of developmental pathways including heart tissue, CNS development, and various aspects of bone maturity.⁴⁷³ BMP receptors are a primary activator of the SMAD family and their associated transforming growth factors. In this contribution, Peterson and co-workers screened a focused library of compounds (7500) for the ability to affect dorsalization of zebrafish embryos from which dorsomorphin 138 was identified. This agent was previously shown to inhibit AMPK activity



Figure 11. Discovery of dorsomorphin.^{473,571}

(K_i between 10 μ M and 20 μ M).⁵⁶⁹ These authors subsequently demonstrated that the AMPK activity of 138 is unrelated to the phenotype of dorsalization and that 138 directly inhibits signaling through the BMP type 1 receptors ALK2, ALK3, and ALK6. A further link was found between BMP signaling and SMAD1/5/8 phosphorylation and ultimately iron-hepcidin homeostasis. A subsequent publication demonstrated the ability of 138 to promote differentiation of embryonic stem cells into cardiomyocytes.⁵⁷⁰ With the target of dorsomorphin established, Peterson and co-workers initiated a program to explore the SAR of dorsomorphin in regard to BMP signaling inhibition.⁵⁷¹ Of particular importance in this study were examinations of the 6-, 3-, and 2-positions of the core pyrazolo[1,5-a]pyrimidine ring. Several synthetic sequences were presented, including the introduction of a diversity element at the 2-position through the choices of appendages at the 3-position of the 1H-pyrazol-5-amine starting material. Reaction of these intermediates with 2-bromomalonaldehyde resulted in the formation of the 6-bromopyrazolo[1,5-a]pyrimidine precursor. Entry of this brominated heterocycle into Suzuki-Miyaura couplings provided another diversity element at the 6-position. Bromination of this heterocyclic core was accomplished via treatment with NBS, and a subsequent coupling provided the final diversity element at the 3-position. From these efforts, DM-3189 (139) was identified as an optimized analog of dorsomorphin (IC₅₀ = 4.9 nM for inhibition of BMP4-induced phosphorylation of SMAD 1/5/8) having good metabolic stability in mouse liver microsomes. These molecules represent important new probes of BMP signaling and may assist the further understanding of the complexities of cellular differentiation.

Molecular Probe: Neuregulin/ErbB4 Signaling Inhibitor (Compound 52).⁴⁷⁴ The past decade has seen several advances in library design and manufacturing. Among



Figure 12. Discovery of neuregulin/ErbB4 signaling inhibitor (compound 52).⁴⁷⁴

these are the establishments of diversity oriented synthesis (DOS) libraries and natural product inspired libraries. DOS libraries are composed of architecturally complex and stereochemically rich small molecules. Such libraries are a stark contrast to the more linear heterocyclic based libraries that dominated screening collections of the past decade. The Schreiber laboratory is the driver of the synthetic technologies behind DOS libraries. In this contribution, Schreiber and co-workers establish a novel Co(I)-catalyzed [2+2+2] cycloaddition between silvl ligated diynes and substituted nitriles in order to advance multifaceted bicyclic pyridines (Figure 12).⁴⁷⁴ Cobalt catalysts have proven useful in several cycloaddition systems and the $CpCo(CO)_2$ catalyst is among the most useful as elevated temperature and irradiation are required to form the active Co(I) species. Selected silyl ligated diynes were reacted with numerous nitriles in the presence of $CpCo(CO)_2$ to provide substituted pyridines. The choice of THF as a solvent provided dramatically enhanced reaction yields and 25 mol % of CpCo(CO)₂ was found to be an optimized catalyst percentage. Interestingly, heat, but not irradiation, was required when using THF for this

transformation. The regiochemistry of the resulting substituted pyridines was confirmed by NOE studies. Treatment of the products from this reaction with TBAF in THF afforded the facile removal of the diisopropylsilyl group. The output was a collection of bicyclic and monocyclic pyridines with various diversity elements interwoven to the structures. A selected number of these agents was screened for inhibition of neuregulin-induced neurite outgrowth via a high-content screen of ErbB4 expressing PC12 cells. One derivative (compound 52: 140) was found to be a potent inhibitor of this pathway. Studies aimed at elucidating the cellular target of 140 are reported to be underway. It is noteworthy that 140 is more reminiscent of traditional library compounds rather than the unique and novel structures found in DOS libraries. This is the case too for (-)-gemmacin (133) discovered from DOS library 5.31 (Figure 9).³⁷⁷

Molecular Probe: Proteinase Activated Receptor-2 (PAR-2) Agonist AC-264613.⁴⁶⁸ The G-protein coupled receptor (GPCR) proteinase activated receptor-2 (PAR-2) plays a critical role in numerous physiological processes, including the inflammatory response. This receptor is activated by proteolytic cleavage of exocyclic peptides on the amino-terminus of the protein by serine proteases (e.g., trypsin). Antagonists of the PAR class of GPCR's have long been sought as a means to circumvent the inflammatory response. Agonists are paradoxically thought to have a beneficial role in smooth muscle relaxation and as cytoprotective agents in the gastric system. Numerous peptide based agonists of PAR-2 have been reported; however, prior to this contribution by Burstein and coworkers⁴⁶⁸ no potent, selective small molecule agonists of PAR-2 had emerged. A HTS campaign of over 250 000 small molecules was performed and two novel structures were found (Figure 13). The first was a relatively elaborate class of hydrazide/hydrazones (141) that displayed good potency and efficacy (6.7 nM and 81% efficacy). The second was a related class of hydrazide/hydrazones with specified chirality (142) that displayed good potency but only modest efficacy (5.2 nM and 30% efficacy). Efforts to expand upon the first chemotype were hindered by poor solubility and synthetic difficulties. Alignment of both lead compounds via a flexible overlay provided good evidence that only modest substituent changes to 142 may provide the necessary efficacy enhancement. Particularly, metasubstitutions on the aryl group attached to the hydrazone were considered important for both potency and efficacy. To explore this hypothesis the authors considered a synthesis utilizing commercially available chiral ethyl ester 144.⁵⁷² Treatment of 144 with hydrazine in ethanol/acetic acid at elevated temperatures in a microwave reactor afforded hydrazide 145 in good yield. A microwaveaccelerated condensation between 145 and the aryl ketone 146 provided AC-264613 (143). SAR exploration confirmed that the *meta*-bromo substitution was essential for both potency (7.5 nM) and efficacy (93%), as the corresponding ortho- and para-bromo substitutions had depressed efficacies (11% and 32%, respectively). The activity of AC-264613 was shown to be highly specific



Figure 13. Discovery of proteinase activated receptor-2 (PAR-2) agonist AC-264613.⁴⁶⁸

for PAR-2 (no activity was noted at PAR-1 or PAR-4) and was confirmed in phosphatidyl inositol hydrolysis and Ca²⁺ mobilization assays. AC-264613 demonstrated efficacy in two divergent in vivo inflammatory assays. AC-264613 represents the first small molecule probe of PAR-2 agonism.

Fluorinating Cleavage of Solid-Phase Linkers.⁴⁰⁸ Fluorine-containing compounds, although not abundantly found in nature, have found their place in modern drug discovery. Introduction of fluorine atoms can dramatically modify the physicochemical, reactivity and biological profiles of organic compounds. Organofluoro chemistry has produced several marketed drugs, including Lipitor, Advair, and Prozac. Recognizing the synthetic limitations to introducing fluorine via high throughput chemistry, Bräse and colleagues devised a solid-phase methodology for the synthesis of gem-difluoro compounds. This is based on the observation that the desulfurative fluorination of dithianes afford gem-difluoro analogs.⁴⁰⁸ Their strategy centered on the immobilization of aldehydes and ketones with a polymer-supported dithiol-based linker 153 to afford dithiane intermediates 14, then employing fluorinating cleavage to release the desired gem-difluoro analogs 148. Commercially available 2-(bromomethyl)acrylic acid 149 was exposed to sodium thioacetate to form dithioacetate intermediate 150 which in turn was anchored onto aminomethyl polystyrene resin 151 under peptide coupling reaction conditions to give resin 152. Since basic hydrolysis would lead to undesired disulfide formation, resin 152 was treated with a solution of HCl in MeOH at 50 °C for 24 h to quantitatively afford the dithiol resin 153. The conversion of 152 to 153 was monitored by gel-phase ¹³C NMR spectroscopy. Resin **153** was condensed with a selection of substituted benzaldehydes and phenylsubstituted methyl ketones 154 in the presence of

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BF₃•Et₂O in chloroform at 40 °C for 24 h generating immobilized dithiane intermediates **147**. Fluorinating cleavage was carried out by treating a suspension of **147** in dichloromethane with *N*-iodosuccinimide (NIS) and HF•pyridine (70%) from -78 °C \rightarrow 0 °C. The desired *gem*-difluoro products **148** were obtained in yields up to 81% starting from resin **153**. NIS was the optimal oxidizing agent since *N*-bromosuccinimide (NBS) or 1,3dibromo-5,5-dimethylhydantoin gave brominated side products.

To determine the robustness and scope of the dithiane linker, solid-supported aniline analog 155 and iodide analog 156 were investigated. Immobilized aniline 155 was successfully acylated with acid chlorides, this was followed by fluorinating cleavage to give gem-difluoro amides 157 in good overall yields (40-60%). Immobilized iodide analog 156 was subjected to various carbon-carbon coupling reactions (e.g., Suzuki, Heck, Horner-Wadsworth-Emmons, olefination, Sonogashira) where it was observed that not only the efficiency of the carbon coupling reactions was low (15-35%) yields) but also the fluorinating cleavage conditions further halogenated products derived from olefin (160) and Sonogashira coupling (161). Immobilized analogs 147 ($R^1 = H$) could be alkylated with n-BuLi/1-bromobutane affording the corresponding pentane analog (147: $R^1 = n$ -Bu; ~16% yield).

Traceless Solid-Phase Synthesis: Thiomorpholin-3ones and 1,4-Diazepan-2-ones.^{333,334} Saruta and Ogiku reported the traceless solid phase synthesis of thiomorpholin-3-ones³³⁴ and 1,4-diazepan-2-ones³³³ via dibenzylation of sulfonium and quaternary ammonium salts, respectively. Thiomorpholin-3-ones are ligands for a wide range of receptors such as 5-HT1B and EP4. There are scant reports on the synthesis of these compounds via solid-phase and the reported methodologies use highly toxic cleavage reagents (e.g., HF) or produce desired products with low purities. A traceless solid-phase approach for the synthesis of thiomorpholin-3-ones was devised where a thiobenzyl linker undergoes intramolecular sulfur alkylation to form benzyl sulfonium intermediates 162 which, upon CsI-mediated debenzylation, releases the desired thiomorpholin-3-ones **163** in a traceless manner (Figure 15). Sulfanyl-ethanols 165 were coupled to Merrifield resin 164 in the presence of DBU to afford thioether-linked alcohols 166. Immobilized alcohols 166 were aminated with N-monosubstituted 2-nitrobenzenesulfonamides 167 under Mitsunobu reaction conditions, followed by removal of the 2-nitrobenzenesulfonyl group with 2-mercaptoethanol. The polymer-supported secondary amines 168 obtained were coupled to chloroacetic acids 169, and the resulting amides 170 underwent intramolecular sulfur-driven cyclization forming benzyl sulfonium intermediates 162. Upon treatment with CsI (dioxane/water (4:1); 95 °C) the benzyl moiety of 162 undergoes an iodine-mediated S_N2 reaction releasing the thiomorpholin-3-ones 163 in high purities (>95%). This methodology offers the advantage that only desired products 163 are afforded, as potential byproducts derived from either S_N2 reaction at the α -position of the carbonyl group or Dithiane linker and fluorinating cleavage



Solid-phase fluorinating cleavage approach



Solid-phase gem-difluoro analog synthesis



Figure 14. Fluorinating cleavage of immobilized dithianes.⁴⁰⁸

 β -elimination of acyclic sulfides remain anchored to the polymer support. Thiomorpholin-3-ones represented by **171** and **172**, where three out of the four diversity sites (R¹ through R⁴) were substituted, afforded products with overall yields of 47–72%. The methodology was extended to construct 7- and 8-membered ring analogs **173** and **174**.

Saruta and Ogiku also developed a traceless solid-phase synthesis approach for biologically active 1,4-diazepin-2-ones **176** (Figure 16).³³³ Analogous to the thiomorpho-lin-3-one synthesis, the traceless route was based on the

Traceless solid-phase sulfonium debenzylation



Figure 15. Traceless solid-phase synthesis of thiomorpholin-3-ones.³³⁴

dibenzylation of quaternary ammonium salts **175** to afford 1,4-diazepin-2-ones **176**. In this approach 4-hydroxymethyl polystyrene resin **177** was aminated with *N*-monosubstituted 2-nitrobenzenesulfonamides **178** under Mitsunobu reaction conditions followed by removal of the 2-nitrobenzenesulfonyl group with 2-mercaptoethanol. The resulting polymer-supported secondary amines **179** were alkylated with alkyl vinyl ketones **180** via Michael addition to afford amino ketones **181** which were then further converted to their corresponding diamines **183** via reductive amination. The polymer-supported diamines **183** were subsequently acylated with chloroacetic acids **169**. Exposure to CsI (dioxane/water (4:1); 95 °C) led to cyclized resin-bound intermediates **175**, which upon final treatment with 2-mercaptoethanol furnished the desired



Figure 16. Traceless solid-phase synthesis of 1,4-diazepin-2-ones.³³³

1,4-diazepin-2-ones **176**. Representative compounds **185–188** were obtained in modest yields (23-48%) and high purity (>91%). The authors demonstrated that substituents at all 4 diversity points were well tolerated in the reaction. The reaction fails with acrolein (**188** R² = H) in place of vinyl ketones (**188** R² = carbon radical).

MCR to 1,4-Diazepin-5-ones.¹⁵⁰ Fenster and co-workers describe the preparation of an extensive library of 1,4diazepin-5-ones as potential tools for the investigation of γ -turn peptidomimetic processes (Figure 17). During the course of the research the authors derived a novel three component synthesis of the diazepinone scaffold. In the first part of the study, diazepinone cores were prepared from piperidone in gram quantities using the ringexpansion protocol developed within their laboratories. Reaction of the *N*-benzyl piperidone **189** with substituted azidoethanols **190a**-**d** gave the iminium ethers **191a**-**d** which were subsequently treated with aqueous bases to give lactams 192a-d. Use of benzylated piperidone allowed for subsequent deprotection $(192a - d \rightarrow 193a - d)$ and diversification of the secondary amine. Various libraries were prepared from the four synthesized diazepinones (193a-d). For example, reductive amination of the four amines (193a–d) with twelve aldehydes generated a 48-member library of N-substituted diazepinones (194). Alternatively the amines 193a-d were treated with various benzoic acids or tosyl chloride to give an 18-member library of amides (195) or the sulfonamides (196) respectively. Further diversification of 195 and 196 was conducted via O-alkylation with a variety of benzyl bromides to produce a 76-member library of compounds (197, 198). A 26-member carbamate library (199) was generated through treatment of compounds 192c and 198a with various isocyanates. In the second part of the study, the susceptibility of the imine intermediate to attack by nucleophiles was exploited. As an extension of previous research conducted by Fenster and colleagues where the electrophilic property of the N-alkyloxazilinium bicyclic ring system (191) was investigated, the authors combined the nucleophilic addition step, in a single-pot fashion, to the ring expansion reaction. In a $2 \times 2 \times 2$ approach, an eight compound library was prepared as a rehearsal using substituted thiophenoxides (R^3 - C_6H_4SH ; 203) as the nucleophile: $201a/b + 202a/b + 203a/b \rightarrow 204a-h$. Initially the eight reactions were performed in a Bohdan Miniblock XT. In an effort to automate the process, the same reactions were then carried out using a Chemspeed SLT100 synthesizer. Comparative results were obtained for the library preparation. Polymeric scavengers were used in the purification of the libraries and the products were isolated with purities $\geq 90\%$. Having established the viability of the chemistry and using four N-substituted piperidones, four aldehydes and eight nucleophiles, a 128member library was generated. Out of the 128 attempted reactions, 117 reactions provided more than 20 mg of product and 113 reactions yielded the products with purities $\geq 85\%$.

Spiroquinolines via One-Pot Multicatalytic and Multicomponent Cascade Reaction.⁴⁰ Multicomponent reactions (MCRs) are highly efficient reactions where three or more reactants can react in a one-pot approach providing a broad range of compound classes that are structurally, electronically, stereochemically, and chemically diverse. These reactions are also very attractive due to their operational simplicity, high atom economy, and high catalytic efficiency. Inspired by the Povarov reaction where activated olefins and aromatic imines react to produce quinolines, Barluenga and co-workers⁴⁰ devised a one-pot multicatalytic-driven multicomponent reaction to synthesize spiroquinolines **210** (Figure 18). One limitation of the Povarov cyclization is the availability of enol ether substrates. This limitation was circumvented in the new cascade reaction by generating enol ethers in situ through catalysis. Thus, a reaction mixture of alkynols 205, aldehydes 206, and anilines 207 was treated with catalytic [PtMe₂(cod)] (cod = cyclooctadiene) and HBF₄. Under these reaction conditions, alkynols 205 suffer Formation and diversification of N-hydroxyethyl-1,4-diazepin-5-ones

R



Figure 17. Synthesis of N-hydroxyethyl-1,4-diazepin-5-one scaffold.¹⁵⁰

intramolecular hydroalkoxylation to form the corresponding exocyclic enol ethers **208**, which in turn, react with in situ generated aromatic imines yielding spiroquinolines **210**. The reaction generally proceeds in high yield (72–89%) giving 1:1 diasteromeric mixtures of spirocyclic products. The cascade reaction appears to have wide scope, accommodating a range of substrate combinations yielding unsubstituted (**211**, **212**), substituted (**213**, **214**), and spirotetrahydro-substituted (**215**, **216**) spiroquinolines. In some instances, single diastereomers are formed (**217**, **218**). When pent-4-yn-1-ol was used, furo[3,2-*c*]quinolines (**219**) were obtained.

Amines Directly from Catalytic Three-Component Ugi Reaction.²⁹⁶ The Ugi four-component reaction has been more widely studied and used than any other MCR.

The process consists of reacting a carbonyl compound, an amine, an isocyanide and a carboxylic acid to yield α -amino acid derivatives **225** (Figure 19). According to the proposed mechanism for this reaction, the addition of the isocyanide to the protonated imine (formed by condensation of the amine with the aldehyde) leads to the nitrilium ion intermediate **224**. Nucleophilic addition of carboxylate to **224** yields an acylimidate derivative, which after an irreversible Mumm rearrangement forms the classical *N*-carbamoyl α -amino acid amide product **225** and water. Researchers at the Max Plank Institute demonstrated that the nitrilium ion **224** can react with the water generated during the imine formation to provide directly α -amino amides **226**. Using benzaldehyde, *p*-anisidine, and *tert*-butyl isocyanide as model substrates, this new Povarov-based multicatalytic reaction



Figure 18. Multicatalytic MCR spiroquinoline synthesis.⁴⁰

three component Ugi reaction is conducted efficiently at 80 °C in toluene in the presence of phenylphosphinic acid (10 mol %). Other catalysts were screened, but phenylphosphinic acid provided the best results in terms of rate of conversion to the desired product. The scope of this reaction was extended using various amines, aldehydes and isocyanides. In all cases reported, the desired products were formed in moderate to good yield demonstrating the broad utility of this new MCR.

Tricyclic Scaffolds via Palladium-Catalyzed 3CR.¹¹⁷ Scientists at Lundbeck identified a novel synthesis of the phenothiazine template **230** using a one-pot reaction consisting of a simultaneous formation of C–S and C–N bonds in the presence of a palladium catalyst (Figure 20). With the use of this new methodology and after optimization of the reaction conditions, the antipsychotics promazine **230a**, chlorpromazine **230b**, flupromazine **230c**, triflupromazine **230d**, and methylpromazine **230e** were all prepared in good yields (50–76%) by condensing 2-bromobenzenethiol, 1-bromo-2-iodobenzene derivatives, and



Path b: new catalytic 3CR catalyzed by phosphinic acid in the absence of R⁴CO₂H

<u>Inputs</u>



Catalysts screened for the Ugi 3CR:



Figure 19. Catalytic three-component Ugi reaction.²⁹⁶

N,*N*-dimethylpropane-1,3-diamine in toluene under microwave or conventional heating conditions in the presence of $Pd_2(dba)_3$ (2.5 mol %), dppf (10 mol %), and sodium *tert*-butoxide (4 equiv). Various benzyl amine and aniline derivatives were also efficient substrates for this novel three component reaction. The utility of this reaction was demonstrated by the one step preparation of promazine **230a** in multigram quantity.



<u>Inputs</u>



Figure 20. Tricyclics via Pd-catalyzed three-component reaction.¹¹⁷

Radical Domino 4CR to Amino Alcohols.¹⁰¹ Clerici and colleagues described a new four component, one-pot procedure for the generation of β -amino alcohols 235 (Figure 21). Previous research performed by the authors demonstrated the utility of a titanium chloride/hydroperoxide system to perform radical versions of the Mannich reaction and Strecker synthesis. A similar titanium III chloride/peroxide system was developed to promote the synthesis of 1,2-amino alcohols via a radical domino multicomponent reaction (e.g., 235a-i). The general applicability of the procedure was established with various aldehydes, and selected primary and secondary amines. The reaction of primary amines with formaldehyde led to a mixture of mono- and disubstituted products (235a,b). When undecamine was used only the disubstituted product was isolated (235c). The authors attribute this to competition between the starting amine and product in the imine formation step. Use of substituted aldehydes, however, resulted in only monosubstituted products. The multiple roles of the titanium and a mechanism for the reaction was proposed. Decomposition of the tert-butylhydroperoxide (234) is induced by oxidation of Ti(III) to Ti(IV).



Figure 21. One-pot, four component synthesis of 1,2-amino alcohols.¹⁰¹

This resulting alkoxy radical (236) then generates the hydroxymethyl radical (237) through hydrogen abstraction from methanol. Simultaneously the Ti(IV) species acts as a Lewis acid in the promotion of the imine formation (238) between the aldehyde and amine. Addition of the hydroxymethanol to the imine generates the radical intermediate (239) which is then reduced by Ti(III) to give the desired products 235.

1,2,4-Oxadiazoles via a Flow Reaction.¹⁶⁷ In a remarkably simple approach, using a single continuous microreactor, Grant and co-workers prepared a series of 1,2,4-oxadiazoles **244** in moderate to good yields (Figure 22). Utilizing an unbroken three-step microreactor sequence, the "in flask" reaction sequence was re-engineered, converting a multistep, multiday synthesis of oxadiazoles into a highly efficient procedure taking <30 min. The utility of the procedure was demonstrated through the use of a variety of aryl and heteroaryl nitriles with various acid chlorides to give products **244a**-**d** in 45–63% yields. When substituted succinic anhydride was used in place of the acid chloride, propanoic acid analogs **247a**-**c** were generated.









Selected products (yield)



Figure 22. Flow reaction to 1,2,4-oxadiazoles.¹⁶⁷

Fluorous Mixture Synthesis of (+)-Cytostatin and Its Stereoisomers.²⁰⁵ Since the introduction of fluorous mixture synthesis (FMS) by the Curran group in 2001, this new solution phase technique has been applied to the synthesis of libraries of drug-like compounds and natural product analogs. One of the most recent examples of FMS is related to the preparation of (+)-cytostatin (248a) and its stereoisomers 248b-d (Figure 23).²⁰⁵ Cytostatin was isolated from the cultured broth of Streptomyces sp.⁵⁷³ This compound is a potent and selective inhibitor of protein phosphatase 2A. It inhibits metastasis of lung melanoma cells in mice and also has potent cytotoxic activity toward leukemia cell lines (IC₅₀ = 42-65 nM). Cytostatin has six chiral centers and its configuration was previously determined by comparison of spectroscopic, physical, and biological properties of a synthetic sample and the natural product. To confirm the structure of cytostatin, the Curran group employed the FMS technique to synthesize two pairs of enantiomerically pure isomers

which were believed to be the most likely candidates for cytostatin. One of the key steps in this work was the coupling of fluorous quasi-racemic mixtures M-252 and M-255. Compounds M-252 each contains three chiral centers and the enantiomeric configurations were encoded by C₆F₁₃ and C₈F₁₇ tags, respectively. Compound M-255 contains two chiral centers and the enantiomeric configurations were encoded by a regular alkyl (zero fluorines) and C₄F₉tag, respectively (Figure 23). Horner-Wadsworth-Emmons reaction of M-252 and M-255, followed by sequential 1,4-reduction using the Stryker reagent and ketone reduction using $LiAl(tBuO)_3H$, generated the sixth chiral center and gave M-257 as a mixture of two pairs of quasi-racemic mixtures, which were then converted to M-258 after several reactions. Since four compounds from two pairs of quasiracemic mixtures of M-258 each has a different fluorine content, they were demixed by the tagbased fluorous HPLC to give four individual fluorous compounds 259a-d. Each of these four separated compounds underwent further transformations followed by fluorous tag cleavage to give four stereoisomers 248a-d. After ¹H NMR comparison and the TLC study of these four compounds and the natural sample, it was concluded that compound 248a has the same configuration as cytostatin and thus confirmed the original structure characterization. This work demonstrated the power of FMS for making natural stereoisomers for structure determination. Since the preparation of four stereoisomers was carried out in one-pot, significant effort was saved in this multistep natural product synthesis.

Fluorous Diastereoselective Synthesis of Cyclic $\beta_{,\beta}$ -Difluorinated α-Amino Acid Derivatives.⁴⁵⁸ Incorporation of fluorine into biologically active compounds to alter drug metabolism and enzyme substrate recognition has become a common practice in medicinal chemistry. One of the applications is to prepare fluorinated amino acid-based peptides and other derivatives. Using a fluorous tag to facilitate product purification, the Fustero group developed a new protocol to synthesize chiral cyclic quaternary α -amino acids for potential biological property studies.⁴⁵⁸ As shown in Figure 24, the fluorous linker was introduced to chiral imidoyl iodide 260 by alkoxycarbonylation to form imino ester (-)-261. After an easy fluorous solid-phase extraction (F-SPE), compound (-)-261 was isolated and subjected to diastereoselective addition of allylzinc bromide generating (-)-262, followed by ring-closing metathesis to afford (-)-263. Transesterification with TBAF in the presence of benzyl bromide cleaved the silylated fluorous tag to yield fluorous chiral cyclic quaternary α -amino ester (-)-**264.** The same research group previously developed similar approaches for both solution-phase and solid-phase synthesis of chiral cyclic quaternary α -amino esters. After side-by-side comparison of these methods with the fluorous method, the authors concluded that fluorous synthesis is more advantageous. It is faster and more cost efficient since it does not require excess reagents to push the reaction to completion as the solid-phase synthesis does. In addition, the fluorous reaction is easy to follow using conventional analytical tools and the product purification by F-SPE is straightforward. Synthetic procedures



Figure 23. Fluorous mixture synthesis of (+)-cytostatin stereoisomers.²⁰⁵

highlighted in Figure 24 have been applied to the preparation of 5-membered ring analogs.¹⁵⁵

Fluorous Synthesis of Structurally Diverse Cyclic Silaketals and Related Z-Alkenes.⁴⁵⁹ Using silaketal to tether allylic or homoallylic alcohols for cross-metathesis is a new strategy for selective synthesis of the Z-isomer of dihydroxyalkenes. This approach has been employed in the target and library synthesis of attenol A, mucocin,

and epothilones. The Nelson group recently reported a fluorous protocol to improve the separation efficiency of unsymmetrical silaketal tethered ring-closing metathesis reactions (Figure 25).⁴⁵⁹ In this work, an allylic alcohol with fluorous-linker **265** was tethered to another allylic alcohol derivative **266** to form silaketal **267**. Ring-closing metathesis of the silaketal generated the desired cyclic product **268** and simultaneously released the fluorous



Figure 24. Fluorous synthesis of chiral cyclic β , β -difluorinated α -amino acid derivative.⁴⁵⁸



Figure 25. Fluorous synthesis of cyclic silaketals.⁴⁵⁹

component **269**. Subsequent ring-closing metathesis of **269** regenerated the fluorous linker **270** as a substituted 2,5-dihydrofuran. The feature of this new protocol is that both silaketal **267** and fluorous linker **270** can be easily purified by F-SPE. Structurally diverse cyclic silaketals **271–274** and related ring-opened diols were prepared by ring-



Figure 26. Fluorous benzaldehyde-based synthesis of 2-arylsubstituted 4-thiazolidinones.⁴⁴⁰

closing metathesis. The same research group has also used a fluorous Grubbs catalyst for the ring-closing metathesis reactions. In this case, both the cleaved fluorous linker and the catalyst were collected in the fluorous fraction after F-SPE.⁴⁵⁷

Microwave-Assisted Fluorous Synthesis of Substituted Thiazolidinones.440 Substituted thiazolidinones possess a wide range of biological activities. Among them, the 2-imine-substituted 4-thiazolidinones have shown selective cytotoxicity to both paclitaxel sensitive and resistant lung cancer cells and are nontoxic to normal human fibroblast.⁵⁷³ The Yan and Zhang groups developed a microwave-assisted fluorous method and synthesized a 2-arylsubstituted 4-thiazolidinone library for quantitative structure activity relationship (QSAR) studies (Figure 26).440 The three-component reaction for the preparation of 4-thiazolidinones 278 involved reaction of fluorous benzaldehydes 277 with excess amounts of the nonfluorous components mercaptoacetic acids 275 and amines 276. Thus, no fluorous starting material was left over and the condensed products containing the fluorous tag were easily isolated from the reaction mixture by F-SPE. The purified compounds 278 were then subjected to Pd-catalyzed fluorous tag cleavage reactions with thiols and boronic acids to form compounds 279a and 279b, respectively. The reaction mixtures were also purified by F-SPE and gave final product purities between 74-100%. To ensure the quality of compounds for cell-based screening, all 60 final products were further purified by semipreparative HPLC. This fluorous tag-facilitated synthetic method has been extended for a small library of the structurally related 2-aryl-substituted 4-thiazinanones.

Table 1. Chemical Libraries Targeting Proteases^a



^a Asterisk is the point of attachment to resin.



• glycogen synthase kinase 3β (GSK- 3β) inhibitors



• dual Akt1/2 inhibitors

• Library 2.1 • Wu [414]

Library 2.5

Castro [78]



 Library 2.2
 Cancilla [72] aurora kinase inhibitor derived from site -specific dynamic combinatorial chemistry

ö

Library 2.6
Murr [279]
DNA-dependent protein kinase inhibitors



• Library 2.3 • Chilin [94] casein kinase 2 (CK2) inhibitors

Library 2.7
Koryakova [214]
GSK-3 inhibitors





 R^2







Mayer [264]
 insulin-like growth factor
receptor (IGF-1R) inhibitors



Table 2. Continued



- dopamine D4 selective ligands
- Library 3.11 Nersesian [282] • histamine H₃ receptor antagonists



• Library 3.12 Sharma [353]
mGluR5 allosteric partial antagonists





 growth hormone secretagogue receptor (GHSR) agonists

 Library 3.14 Wijtmans [409]
 histamine H₃
 receptor agonists



R

Library 4.3

• Gitto [165]

in DBA/2 mice

anticonvulsant activity

against audiogenic seizures

'N

 \dot{R}^2

• Library 4.4

• Domling [134]

Bcl-2 family protein-protein disruptor

Library 4.1

• Bench [47]

activators of

neurite outgrowth

Library 4.2

• Kaminski [207]

(MES) and/or pentylenetetrazole induced seizure (scPTZ) models

Table 4. Continued



Table 5. Chemical Libraries Yielding Cytotoxic and Antiinfective Agents^a



Table 5. Continued



^a Asterisk is the point of attachment to resin.

Table 6. Selected Molecular Probes

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- SMN2 promoter activator
 - TbcatB inhibitor

Table 6. Continued







sequential palladium-catalyzed coupling reactions



 Severinsen [341] multistep sequence using safety catch linker

Solution-phase

Ar

Alacid [13]

· Heck reaction of acrolein

catalyzed by Kaiser oxime

resin derived palladacycle

diethyl acetal with aryl halides



 Alaimo [14] of ArNO₂, RCHO and Danishefsky's diene or silyl ketene acetal



H₂N_×

0=s

Gutierrez [177]

R¹.NH

• CITi(O/Pr)₃-promoted reductive amination, sulfonylation of the resin-bound amine, Suzuki

ΝH₂

cross-coupling, and acid-mediated cleavage

_Ns

resin-bound secondary amine

Chang [84]

• Hahn [178]

multistep sequence

alkylation of
resin-bound benzyl sulfinate
with allylbromide, ozonolysis

and traceless sulfinate elimination

ΝO2



R¹

ö

diketene and water

H₂N







• Choi [97] cleavage of Marshall resin-bound caboxylic acids with NH₂OH



 Wiehn [408]
 fluorinating cleavage
 of resin-bound dithianes with NIS/HF



- Fontaine [153] 3CR of R¹CHO, R²NH₂, and TMSCN then IBX oxidation
 - ArN2⁺ TsO⁻

 Filimonov [151] treatment of ArNH₂ with polymer-supported nitrite

from tropinone

Nicolaou [284]



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m-, and p-substituted iodobenzoic acid derivatives • Pirali [302] • 4CR of diamines, R³CHO, R²NCO, and R¹CONHCH₂COOH



Prokopcova [310]
MW-assisted one-pot diboration/Suzuki cross -couplings of alkynes



In-catalyzed domino reaction of ArNO₂, RCHO and Danishefsky's diene or silyl ketene acetal

 from aryInitriles and activated carbonyls in a single continuous microreactor sequence condensation of isocyanoacetate and ArN=C=S using microfluidic reaction chips and packed immobilized-reagent columns

 condensation of thiazolidine and β-enaminonitrile

derivatives



azide-alkyne cycloaddition (RuAAC)

presence of Lawesson's

reagent

as ammonia source



Table 8. Continued

Boc \NH Ą $R^2 \rightarrow$

R¹

Farran [148]
 treatment of bis-Boc
protected 2,5-diketo

-piperazines with LiHMDS and RCH₂Br

`N

Ŕ2

• Alizadeh [19] • 4CR of R¹NH₂, R²NH₂, diketene, and nitrostyrene

R¹



• Waldo [398] iodocyclization of O-methyloximes of 2-alkyn -1-ones then various Pd-catalyzed reactions



 Volonterio [397] volonterto [397]
 condensation of carbodiimides with malonic acid monoethylesters, cyclization of the resulting *N*-acylureas, and the C-alkylation with R⁴X



 Yadav [422]
 3CR of carbonyl compounds, • 3CR of RCHO, homoallylic alcohols, and homoallylic alcohols, and nitriles trimethy silyl azide

Yadav [423]
3CR of acyclic or cyclic active methylene ketones

Boc



• Xiao [419] • [PdCl₂(CH₃CN)₂]-catalyzed 3CR Michael addition/cyclization anions, vinamidinium /cross-coupling of various 2 -(1-alkynyl)-2-alken-1-ones with MeOH and allyl chloride chloride, and NH₄OAc



• Zhang [436] • 3CR of DMAD, R¹NH₂, and R²CHO



Asghari [30]
3CR of ethyl acetamido -cyanoacetate, dialkyl acetylenedicarboxylate, and Ph₃P



Akbas [11]
Biginelli 3CR of β-diketone, RCHO, and thiourea



 Adamo [4]
 3CR of 3,5-dimethyl 4-nitroisoxazole, ethyl 2-chloroacetoacetate and ArCHO



Ackermann [3]
 Cu-catalyzed MCR of alkyne, NaN₃ and R²X, followed by



• Liu [248] • 3CR of acid chlorides, terminal



R³ Noth [287]
3CR of maleimides,

aldehydes, and amines

R⁵ 'nн ,н ∱R⁴ R³

• Elders [140] • 3CR of amines, aldehydes/ketones α-acidic isocvano amides or esters

• Alizadeh [20] • 4CR of R¹NH₂, diketene, R²NH₂ and dibenzoylacetylene

 \mathbb{R}^2

R¹ нŃ

Bn

Bn



R²



 \dot{R}^2

R¹



Alizadeh [17]
5CR of 1,n-diamines, diketene, dibenzoylacetylene in the presence of Ph₃P





multistep sequence

triazole arylation









• Dou [135] • 3CR of 1,3-diketones, aldehydes, and amines induced by TiCl₄/Sm

Brown [65]
 multistep sequence



R³

 \mathbb{R}^2

'N

• Lee [232]

'R

∠R²

Ĭ

Reddy [316]
3CR of allylsilanes,

R¹CHO and R²CN

ΗÑ.

่่่≈ุ่∧

Boo ́юн бн • Krishna [222] $\begin{array}{l} \text{from } L = \text{Amino acids via} \\ \text{diastereoselective Baylis-Hillman} \\ \text{reaction of } \textit{N-allyl-Boc } \alpha \text{-aminal,} \\ \text{followed by RCM and dihydroxylation} \end{array}$



Bremner [61]
 one-pot domino aldehyde
/amine condensation, [3,3]
-aza-Claisen rearrangement,
imine-allene cyclization of
proportubrities and PCHO



R³

R

acetylenedicarboxylate

• Alizadeh [16] • 3CR of RNH₂,

ArSO₂NCO, and dialkyl

• Khalili [210] 3CR of β-dicarbonyl compounds with

arylglyoxals in the presence of NH₄OAc in water

R¹



• Bonger [58] • tandem Staudinger/aza -Wittig/Ugi 3CR on a L-ribose -derived 4-azido aldehyde

-R²

o.

`Bn



• Kralj [215] • reaction of *N*-protected α-enamino lactams derived from 2-pyrrolidinone with RNHNH₂



Li [240]
 4CR of β-aroythioamides, aldehydes, acetonitrile derivatives, and 1,4-dihydropyridines followed by S-alkylation

Table 8. Continued



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Hioki [185]

oxidative release from azomethine linkage on a solid-support

by 2-aminobenzylamine or 2-aminobenzamide with air

Solution-phase

 \mathbb{R}^2

R¹

 \mathbb{R}^2

and isocyanides

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°0

·NH

R

N^{-R¹}

R²



Xiao [418]
Au-catalyzed 3CR of aldehydes, amines, and alkynes

Shih [357]multistep sequence from N-substituted methyl ester serine

Vieira [396]
Rh-catalyzed 3CR of 2-vinylbenzaldehydes, ArNH₂ and CO



• Yang [425] • 3CR of ArNH₂, ArCOCI, and heteroaromatic amino ester





• Zuo [15] • MW-assisted 3CR of 2-chlorophenols, chloro -acetylchloride, and R²NH₂



 Zhang [435] • Cu-catalyzed 3CR and Diels-Alder reaction

• Frankowski [154] • tandem Diels-Alder /Schmidt reaction then skeletal diversification



 Adib [5]
 3CR of 2-aminohetero
-cycle, ArCHO, and
imidazoline-2,4,5-trione • El Kaim [139] • o-iodonitrophenol in Ugi-Smiles reaction coupled with Heck cyclization



isocyanides

reaction with aryl iodides



Akai [10]
 Akai [10]
 CR of 3-TBDMS-benzynes
 Additional derivatives then
Hiyama cross-coupling
 with and indides



 \bullet Almansa [25] \bullet 3CR of R $^1COCH_2R^2,$ $R^3CHO,$ and 3-aminopyrazole

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Table 9. Continued

• Bouillon [59] • tandem condensation of R^1NH_2 , 2-NO₂-Ph R^2SO_2CI , and BrCH₂COR³



 Karapetyan [209]
 domino reaction via ZnCl₂-catalyzed conjugate addition/cyclization sequence of 1,2-diaza-1,3-butadienes and 1,3-bis(silvlenol ethers)



 Li [244]
 MW-assisted reaction of substituted methyl anthranilate, and iso(thio)cyanates in DMSO /H₂O without catalyst or base



Bogolubsky [53]
 cyclization of 2-amino
 -thiophen-3,5-dicarboxylates

with RCN then hydrolysis

0=

N-DNA H

Buller [67]
DNA-encoded library via Diels-Alder cycloaddition

 Jainta [191]
 MW-assisted condensation
 of unprotected amino acids
 by a phosphite-promoted one-step coupling reaction





Ar

CN



of phenols



 Kim [211]
 Cu-catalyzed 3CR
 of alkynes, ArSO₂N₃,
 NH₄CI then intramolecular N-arylation

 R^1

• Li [239] • multistep sequence via sequential C-H functionalization

R¹

Dahl [117]
 Pd-catalyzed, MW-assisted
 3CR of 2-Br-thiophenol, amine,
 and 1-Br-2-iodobenzene



Liu [251]
 DOS via thiourea/AuCl₃ /AgOTf-catalyzed annulations of aryl epoxides



• Gao [158] • [3+2] cycloaddition of 5-benzenesulfonyl -3,4-dihydro-1*H*-pyridin-2-one derivatives with azides or isocyanides

 R^2 N R^1 0

Garcia-Cuadrado [159]
DOS utilizing a Petasis 3CR followed by a tandem aza-Cope-Mannich cyclization



• Ding [132] • tandem addition-cyclization of 2-alkynylbenzenamines with RNCS catalyzed by AgOTf



Balazs [34]
MW-assisted multistep sequence from cyclic β-amino acids

Ċh7



Banfi [37]
Ugi/Mitsunobu reaction of α -hydroxybenzoic acids, glycolaldehyde dimer $R^2 N H_2$ and $R^3 C N$



• Cho [95] multistep sequence



sequence

R

Matsuya [262]
3CR and cyclization of

N-tosylimines and TMS -substituted propiolate mediated by DABCO

R²

Salcedo [331]
Ugi-4CR/postfunctionalization

 Carpenter [76]
 multistep sequence
from 2-OH-PhCOMe and 4-ketopiperidine



R²

`R³

Mao [260]
allenyl-hydroxy ester allylated or propargylated

then subjected to Rh(I)-catalyzed carbocyclization

·R'

F₃C

0=

 $-R^1$

• De Silva [126] • MW-assisted 4CR of (2-NO₂)ArCHO, cinnamic acids, R³NC, and R²NH₂ with Fe/NH₄Cl

• Coldham [103]

group

3CR of an amino-acid or amino-ester or hydroxyl-

amine, an alkene or alkyne dipolarophile and an aldehyde

bearing a halide as a leaving



• Cao [73]

Pd-catalyzed
 Sonogashira

coupling-carbonylation -hydroamination sequence

in phosphonium salt-based ionic liquid

R¹

-P'



• Miyazaki [273]

intramolecular cyclization of o-amino -arylalkynes via Au(III)

polymer supported catalyst





 Milinkevich [270]
 condensation of 3-chloro
pentane-2,4-dione with thio
-amides and subsequent Wittig
olefination, followed by nitrile oxide 1,3-dipolar cycliaddition

-R





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Table 9. Continued



Table 10. Continued



Mont [274]
 multistep MW
-assisted DOS

 Corres [109]
 sequential Ugi reaction of ArCHO, 2-acetyl or benzoylanilines, cyclohexyl or benzyl isocyanides, and 2-azidobenzoic acid, followed by a Staudinger/aza-Wittig cyclication with Pho-P cyclization with Ph3P



• Bonfield [57] • 6CR of ArNH₂, HCOH and phenylacetylene

Sayyafi [336]
 3CR of phthalhydrazide, dimedone, and aromatic aldehydes



Shakibaei [351]
MW-assisted 3CR of 6-quinolinol, ArCHO, and urea



An [26]
 DOS of benzopyranyl heterocycles from s-cis-enones



• condensation of 2-NO₂PhCHO with 2-aminobenzylalcohol



 Barluenga [40]
 one-pot multicatalytic and multicomponent cascade reaction of alkynes, RCHO, ArNH₂



Butler [68]
condensation of 2-NO₂PhCHO with 2-aminobenzylalcohol



Jakubec [192]
nitro-Mannich/lactamization cascade of y-nitro esters with cyclic imines



 Lai [225]
 Yb(OTf)₃-catalyzed 4CR of various anisidines, ArCHO, isobutyraldehyde, and 4-hydroxycoumarins/dimedone



Li [243]DOS via Pd-thiourea -catalyzed alkoxy -carbonylative annulation of 1,2-hydroxymethylaryls





• Ding [131] • 3CR of thiazoles or benzothiazole carbenes, disubstituted ketenes and activated alkynes



Barluenga [41]
 one-pot multicomponent cascade reaction of alkynes, ArCHO, RNH₂ catalyzed by Pd-complexes



• Beaumont [46] • 3CR of indole-derived oxo acids, $R^1 N H_2$ and $R^2 N C$



Christodoulou [98] multistep sequence



Litvinov [247]
3CR of substituted isatins, cyanoacetic acid derivatives, and carbonyl compounds or phenols



Shi [356]multistep sequence from 2-CI-nitro -heteroaromatics



Duan [136]
3CR of alkynes, Fischer carbene complexes, and benzaldehyde hydrazones

and active methylene

compounds





Sml2-mediated linker cleavage-cyclization in traceless fluorous synthesis



Sunnemann [373]
multistep DOS



Parenty [297]
quaternarization of *N*-hetero -cycle, followed by a cascade reaction involving nucleophilic addition, substitution, rearrangement, and oxidation



Ohta [290]
Cu-catalyzed domino 3CR coupling-indole formation-N -arylation under microwave irradiation

Table 10. Continued







 various synthetic applications of aryl tetraflates



synthesis

C₈F₁₇



C₈F₁₇

C₈F₁₇



Miura [444]
 catalytic enantioselective cyclopropanation of allylic alcohols

Ts



Hensle [445]
Grubbs-Hoveyda-type metathesis catalyst

Table 12. Continued



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