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Comprehensive Survey of Chemical Libraries for Drug Discovery and Chemical Biology: 2009

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This is the thirteenth installment of the comprehensive survey series in high throughput chemistry.¹ Biologically active libraries reported in 2009 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 370 libraries and 24 molecular probes extracted from 355 literature citations.^{2–491} Approximately 90% of the citations originated from academic laboratories, with the majority of these from European and Asian laboratories. Solution-phase methodology accounted for *ca*. 85% of chemical library synthesis.

This year's 18 vignettes include (a) biologically active libraries: HDAC1/HDAC2 inhibitors,¹³⁴ H₃ antagonists,¹³⁶ glucagon receptor antagonists,¹⁷⁹ purinergic P2Y₁₂ receptor antagonists,²²¹ heat shock protein 90 (Hsp90) inhibitors,⁴¹ selective norepinephrine reuptake inhibitors,⁷³ allosteric modulators of mGluR4²⁸⁷ and mGluR5⁷¹ Bcl-2 inhibitors via DOS library;¹⁸² (b) molecular probes: Ned-19³⁴⁰ and DG-041;³³⁸ (c) high throughput chemical methodology: catch and release synthesis of substituted guanidines.²⁸⁹ and substituted pyrimidines via a 3-component reaction;¹⁵⁸ and (d) fluorous technology: displaceable fluorous dihydropyran³²² and isonnitrile linkers,³²⁷ fluorous synthesis of 1,4benzodiazepine-2,5-dione, ³²⁷ piperazinedion-fulsed tricyclic³²⁸ compound libraries and a fluorous mixture synthesis of natural product resorcyclic acid lactone library.³²⁹

Related publications and reviews appeared in 2009 on microwave-assisted of *N*-heterocyclic synthesis³⁵⁶ and convertible isonitriles,³⁶⁵ click chemistry,³⁵⁸ dynamic combinatorial

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chemistry,³⁶³ large scale preparation of silicon-functionalized SynPhase lanterns,³⁷⁰ kinase inhibitors,³⁵⁷ heat shock protein 90 inhibitors,³⁷⁶ biologically active benzoannelated nitrogen heterocycles,³⁶¹ library automation and analysis,^{360,368} library design,³⁷⁵ NMRased screening,^{377,378} fragement libraries and fragement hopping,^{366,367,369} review of Ellman's libraries,³⁷⁴ solid-phase resin specifications,³⁵⁹ microelectrode arrays for monitoring ligand-receptor binding events,³⁶⁴ yactoliter-scale DNA reactor for small molecule evolution,³⁶² DNA encoded libraries,³⁷⁹ and fluorous chemistry and separations.^{380–383}

Selective HDAC1/HDAC2 Inhibitors.¹³⁴

Histone deacetylases (HDACs) are enzymes involved in the remodeling of chromatin. Several classes of HDAC inhibitors have been found to have potent and specific anticancer activities and the common pharmacophore identified contains a surface recognition domain, a linker region and a metal binding region. Kattar and co-workers set out to improve upon the selectivity and tolerability of Zolinza 1, Merck's first-in-class mixed HDAC 1, 2, 3 and 6 inhibitor used for the treatment of cutaneous manifestation of T-cell lymphoma (Figure 1).¹³⁴ In their initial library, keeping the metal binding domain constant and varying both linker and surface recognition domains, scaffold A had already been identified. Reacting 4formylmethyl-benzoate and polystyrene bound sodium triacetoxyborohydride with 48 R¹amines, followed by hydroxamate formation with NH₂OH for their first follow-up library, 3chlorobenzylamine was found to be preferred in the R¹-position over phenethyl or aniline analogs, yielding compound 2 (HDAC IC₅₀ = 348 nM). Modifying the metal binding domain next, 2-aminophenols were chosen to replace the hydroxamate moiety. R¹substituted 2-nitrophenols 3 were reacted with bromo-Wang resin, cesium carbonate and DMF under microwave irradiation at 40 °C. The nitro group was reduced with SnCl₂ in DMF and the resulting resin-bound anilines 4 coupled with the linker unit 4-chloromethyl benzovl chloride 5, DMAP and DIEA in DCM. Attachment of R²-amines was accomplished using sodium iodide and a proton sponge in DMF. Final compounds $\mathbf{6}$ were released from resin with TFA/DCM. Aromatic substituents in R¹ and anilines in R² demonstrated 10-30fold increased potency for HDAC1 (e.g. 7, $IC_{50} = 10 \text{ nM}$, cellular assay = 1,735 nM). To improve cellular potency the aminophenol metal binding site was replaced with a primary amine. Mono protected R¹-substituted di-anilines were bound to aldehyde resin 8 via reductive amination utilizing sodium triacetoxyborohydride in AcOH/DCE. 4-Chloromethyl benzoyl chloride was attached next, followed by reaction with surface binding domain R²amines, sodium iodide and a proton sponge. Exposure to TFA/DCM yielded final products 9. Amino spirocyclic building blocks in the R²-position retained affinity towards HDAC1 and improved selectivity over HDAC3 as well as improved potency in the cellular assay. Unfortunately this series exhibited activity against hERG (10: HDAC1 IC₅₀ = 4 nM, cellular assay = 82 nM, HDAC3 IC₅₀ = 9,617 nM, hERG IC₅₀ = 1,597 nM). Eliminating one aminofunctionality by switching to a terephthalamide linkage successfully mitigated hERG liabilities. This last set of compounds was synthesized via solution-phase, starting from 11 and mono protected R¹-substituted di-anilines which were combined with PS-CDI and HOBt in DMF. After ester saponification, the spirocyclic amine was coupled using PS-CDI again. The final 26 compounds 12 were obtained after deprotection with TFA/DCM.

Compound **13** showed the best overall profile (HDAC1: $IC_{50} = 8$ nM, cellular assay = 103 nM, HDAC3: $IC_{50} = 5,102$ nM, hERG $IC_{50} = 12,390$ nM) and demonstrated *in vivo* efficacy in an acute PD HCT-116 xenograph model study. This successful combination of solid and solution phase library synthesis led to the efficient optimization of a novel HDAC inhibitor series, continuously improving additional properties within each library iteration.

H₃ Antagonists.¹³⁶

The histamine H₃ receptor is an attractive G protein–coupled receptor drug target that regulates neurotransmission in the central nervous system. There has been considerable effort by both academic and industrial laboratories to develop potent and selective H_3 receptor antagonists for the potential treatment of attention-deficit hyperactivity disorder, dementias, schizophrenia, as well as obesity and sleep disorders resulting in a refined H₃ antagonist pharmacophore model containing two basic nitrogens separated by a spacer and a central core that also carries a polar group and a lipophilic residue (Figure 2). After having completed its total synthesis, Kennedy and co-workers recognized that the marine alkaloid dispyrin 14 perfectly maps onto this pharmacophore model.¹³⁶ Dispyrin 14 indeed displayed some H₃ antagonist activity (IC₅₀ = 2.35 μ M) and was used as the starting point for a natural product guided iterative parallel synthesis campaign. Starting from 3-bromo-4methoxyphenylethylamine 15 five heterocyclic carboxylic acids R¹ were coupled using DIC, HOBt and DIEA in DCM. The methyl ether was removed with BBr3 and the resulting phenols alkylated with five R²-aminoalkyl chlorides under microwave irradiation conditions generating 25 different compounds 16. All the compounds showed activity against H₃ with more potent compounds containing an ethyl pyrrolidinyl residue in the R² position (K_i 's < 200 nM), with the best compound 17 (H₃ K_i = 80 nM, H₃ IC₅₀ = 180 nM). Retaining the 4bromo-thiophene residue from 17, functionalized pyrrolidines were explored next. Phenol 18 was reacted first with 2-bromo-1,1-dimethoxy ethane and then with tosylic acid to yield aldehyde 19. Final compounds 20 were obtained via reductive amination utilizing resinbound triacetoxyborohydride together with the functionalized pyrrolidines. No potency improvement was observed in final compounds 20.

The last library reexamined the R¹ residue, including additional heterocycles and aromatic moieties. Protecting bromomethoxy phenethyl amine **21** as the phthalimide by reacting it with 1,2-dicarboxybenzene, DIC, HOBt and DIEA, followed by BBr₃ demethylation yielded phenol **22**. Alkylation with chloroethyl pyrrolidine and deprotection with hydrazine, both under microwave conditions, followed by coupling with a diverse set of R¹-acids with DIC, HOBt yielded final compounds **23**. Five membered heterocycles in the R¹-position showed superior activities compared to pyridine and substituted benzenes, with 5-oxazole (**24**) and 2-thiazole (**25**) displaying the best potencies, $K_{is} = 32$ nM and IC₅₀ = 72–83 nM, respectively. This iterative parallel library campaign successfully optimized H₃ antagonism properties of a natural product lead structure over 30-fold.

Human Glucagon Receptor Antagonists.¹⁷⁹

The glucagon receptor is a member of the class B G-protein coupled family of receptors. Glucagon maintains glucose homeostasis during the fasting state by promoting hepatic

gluconeogenesis and glycogenolysis. Antagonizing the glucagon receptor is expected to result in reduced hepatic glucose overproduction, leading to overall glycemic control and a possible treatment for type 2 diabetes. Only a few classes of non-peptidic glucagon receptor antagonists are known (Figure 3). Madsen and co-workers had previously described βalanine and isoserine urea based human glucagon receptor (hGluR) antagonists 26 and 27, whereby 27 showed improved selectivity over the related human glucose-dependent insulinotropic receptor (hGIPR). Here the authors explored the replacement of the urea linkage with a heterocyclic scaffold, thus rigidifying the molecule.¹⁷⁹ A library assisted optimization strategy was pursued, combining solid and solution phase synthesis approaches, with the goal of finding a novel potent, orally available, hGluR selective antagonists. For the first library on solid support, 4-formylbenzoic acid was coupled to deprotected Fmoc- β -Ala-Wang resin with HOBt/DIC, followed by reductive amination with R¹-amines and treated with Fmoc-NCS to afford resin-bound Fmoc protected thioureas. Thiazole formation was accomplished through reaction with a-bromoketones after Fmocdeprotection and final compounds 28 were obtained after cleavage from resin with TFA. Over 800 analogs were prepared via this procedure. Compounds were also prepared via solution-phase starting from 4-formylbenzoic acid methyl ester 29, which was reductively aminated with R¹-amines utilizing sodium cyanoborohydride. The benzylic amines **30** were converted to their corresponding thioureas employing different methods depending on the reactivity of the respective secondary amine. The aminothiazole 31 was obtained via reaction with α -bromoketones in acetic acid. The methyl ester was cleaved with NaOH and the resulting carboxylic acid **32** was coupled with either β -alanine or (*R*)-isoserine methyl ester, followed by hydrolysis to yield the final compounds 33 and 34, respectively. Compounds **35** with the aminothiazole directly attached to the aromatic ring were prepared in similar fashion both on solid support and solution-phase. 4-Nitrobenzoyl chloride was reacted with resin-bound β -alanine, the nitro group was reduced with SnCl₂, followed by reductive alkylation with R¹-aldehydes. Thiourea **36** was obtained after reaction with Fmocisothiocyanate. After deprotection, cyclization with a-bromoketones and cleavage from resin final compounds 35 were obtained in high yields and purities. Using a solution-phase approach, compounds from this type were obtained by reductive amination of 4aminomethyl benzoate 37 with R¹-aldehydes, followed by reactions with EtO₂C-NCS, NaOH, α -bromoketones and subsequent EDAC, HOBt coupling with β -alanine methyl ester and hydrolysis to give 38. Finally, compounds 39 containing a 5-alkylthiazole core were prepared starting with a Knoevenagel condensation of 4-methyl benzoate 40 with phenyl acetonitriles, followed by reduction with NaBH4 in THF and treatment with dithiophosphoric acid O,O-diethylester. The resulting saturated thioamide 41 was reacted with α -bromoketones and coupled to β -alanine methyl ester followed by hydrolysis. Aliphatic R¹ groups were not well tolerated. Compounds with 4-CF₃-, 4-OCF₃- and 4-SCF₃phenyl in the R¹ position displayed good binding affinities and superior rat PK properties. Compound 42 (hGluR IC₅₀ = 93 nM, hGIPR IC₅₀ = 1,100 nM) showed high oral bioavailability (58%), low clearance (1mL/min)/kg, long plasma T_{1/2} (228 min after i.v. administration), and extremely high plasma exposure ($C_{max} = 2100 \text{ ng/mL}$) in the rat (3 mg/kg, p.o.). Changing the β -alanine to isoserine in this series did lead to a loss of mGluR activity. Compounds with modified core structures retained binding affinities but displayed less favorable PK profiles. The SAR for R² and R³ residues is of minor importance as long

as the substituents on R² are lipophilic and in *meta* or *para* position of the phenyl ring. Compound **42** was tested in a non-human primate model of hyperglucagonaemia and hyperglycaemia. It dose-dependently decreased glucagon stimulated glycaemia and abolished the hyperglycemic effect of exogenously administered glucagon completely at i.v. doses of 1 and 3 mg/kg. Compound **42** also showed high plasma exposure and a long plasma half-life in monkeys. This library approach demonstrated that the urea linkage in previously reported hGluR antagonists can be successfully replaced by a number of different thiazole cores and established SARs for binding, selectivity and PK properties for this novel chemical class.

Purinergic P2Y₁₂ receptor antagonists.²²¹

Plavix® (clopidogrel, 43) is an antiplatelet agent approved for stroke and myocardial infarction in patients with atherosclerosis. Its mechanism of action is thought to proceed via metabolism to acid 44 followed by irreversible inactivation of P2Y₁₂, a platelet specific GPCR (Figure 4). This in turn leads to a reduction in adenosine diphosphate (ADP)stimulated platelet aggregation and the formation of platelet aggregates thereby providing therapeutic benefit. Because 43 is a prodrug requiring metabolic activation, it cannot be used effectively in patients that require emergency treatment (delayed time of action) or in patients who cannot metabolize 43. Recognizing the shortcomings of 43, Parlow and colleagues sought to find a new class of P2Y12 receptor antagonists having direct action on the receptor.²²¹ P2Y₁₂-based quinoline antagonists were previously reported by Berlex and this class was used as a starting point for library design and SAR exploration. Initial efforts focused on replacing the quinoline moiety employing polymer-assisted solution-phase chemistry. Piperazine derivatives 46 were coupled with heteroaromatic carboxylic acids 47 in the presence of resin-bound carbodiimide and HOBt. Excess reactants were sequestered with resin-bound isocyanate and a resin-bound secondary amine providing clean intermediates that, post TFA treatment, afforded desired carboxylic-acid-containing analogs 48. Hundreds of compounds were prepared using this approach. The library was evaluated in a human-derived P2Y₁₂ receptor binding assay, screening for % inhibition at 10 μ M. K_i values and inhibitory effects in a human platelet rich plasma (PRP) assay were obtained for the more potent compounds. The results from the first round of library evaluation indicated that replacing the original quinoline group with pyridine generally produced compounds with nanomolar K_i values, although with poor PRP inhibitory effects. Since the most potent compounds had one or two phenyl groups on the pyridine ring (49 and 50, respectively), the authors selected 49 as the lead for further SAR exploration and optimization. The next step consisted of SAR exploration around the piperazine moiety **51**. A broad range of alkylating, acylating and sulfonating agents were used in conjunction with a scavenger resin to remove residual electrophile reactants. Biological assays showed that activity was driven by the carbamate group where the longer the aliphatic carbamate chain, the greater receptor affinity binding and, importantly, inhibitory PRP activity (53-57: $K_{is} = 2440$ to 11 nM; $IC_{50}s = >100$ to 15 µM).

The next SAR campaign focused on the 4-position of the pyridine ring. The authors opted to make chloro- and phenol-containing intermediates (**58-60**) to use these functional groups as handles to introduce chemical diversity. Phenol **58** was *O*-alkylated with alcohols under

Mitsunobu reaction conditions (DEAD, PPh₃, THF) or with halides (Cs₂CO₃, KI, DMF) ultimately yielding carboxylic acid ethers **61-66**. Although the SAR was rather flat, a trend emerged for increased PRP activity with the phenolic ethers (**65**, **66**: IC₅₀ = 3.8 and 1.9 μ M, respectively).

In addition to 4-ether substitution, 4-amino substitution was explored. This was carried out by treating carbamate **55** with 20 equiv of amine to displace the 4-chloro group yielding 4aminopyridine ethyl carbamates **67-76**. Simultaneous variation of the 4-position and the carbamate moiety was achieved using the same 4-chloro displacement strategy with **60** followed by alloc deprotection, *N*-piperidine carbamoylation and TFA deprotection. Small secondary alkyl amines were preferred over tertiary amines (IC₅₀s for **67** and **68** versus **69**) while amino ethers exhibited higher PRP inhibitory activity (**70**, **71**); particularly effective were the amino piperidyl and amino piperazine groups (**72-75**). Lastly, increasing the length of the alkyl carbamate chain led to further increases in K_i and IC₅₀ (**76**: $K_i = 7$ nM; IC₅₀ = 0.77 μ M). Evaluation of the most potent candidates in this series led to the discovery of **77** with a satisfactory *in vivo* profile, oral bioavailability, and some 340-fold more selective for the P2Y₁₂ receptor versus its closest homolog, the P2Y₁₃ receptor.

Heat shock protein 90 (Hsp90) Inhibitors.⁴¹

Hsp90 has gained attention in the pharmaceutical industry due to its participation in multiple cell signaling pathways in cancer cells (e.g., PI3K/Akt). Hsp90 is an ATPase protein that acts as a chaperone, binding to multiple oncology relevant client proteins (e.g., Her2, cKit, MET), stabilizing these proteins so to permit their cell signaling function. Inhibition of Hsp90 prevents the necessary folding required to bind and stabilize client proteins. This results in the degradation of the client proteins via the ubiquitin proteasome pathway making Hsp90 an attractive oncology target. Geldanamycin and close synthetic analog 17allylamino-17-demethoxygeldanamycin (17-AAG) bind to the ATP active site in the Nterminal domain inhibiting Hsp90 function validating this to be a viable mode of action to disrupt Hsp90's biological function. Poor water solubility and hepatotoxicity limit the clinical utility of these agents and thus, new small molecules are sought targeting the ATP binding site to inhibit Hsp90 function. Radicicol is an inhibitor of Hsp90 via ATP competition binding. Promising results have been noted by several research programs with resorcinol-based Hsp90 inhibitors. However, a common limitation to this compound class is that the phenol groups can be substrate sites of metabolic degradation processes (e.g., glucuronidation) resulting in fast in vivo clearance.

Cho-Schultz and colleagues focused their attention on compound **78**, an amide-containing polyphenol and ATP binding site-directed inhibitor of Hsp90 (Figure 5).⁴¹ A two-stage solution-phase strategy was devised to explore the chemical space and develop an SAR around **78** to produce new potent Hsp90 inhibitors without phenol groups. The first library of compounds focused on the solution-phase synthesis of **79**. The objective was to improve binding affinity and reduce or eliminate glucuronidation by increasing hydrophilicity via the introduction of basic amino fragments. Library synthesis commenced by the addition of Grignard intermediate **81**, derived from *m*- and *p*-1,3-dioxane-subsituted bromobenzene to *N*-vinyl pyrrolidinone **82** affording 2-pyrrolines **83** (23–29% yields). These cyclic imines

were reduced to 84 with NaBH₄ and then coupled to phenol-protected benzoic acid 85. Deprotection of the phenol groups with either TFA or HCl resulted in aldehydes 86 that were subjected to reductive amination with ca. 90 amines to afford the 178-compound library 79. Compounds were tested in a tritium-labeled-ligand competitive binding assay (K_i) followed by the measurement of Akt-degradation in H1299 lung cancer cells (IC₅₀). The SAR indicated that para substituted compounds had higher binding affinity against Hsp90 and demonstrated superior activity in the cell-based assay ($K_{is} = 5.4-40$ nM; IC₅₀s = 112–668 nM) versus the *meta* substituted compounds ($K_{is} = 60-100$ nM; IC₅₀s = 2100– 4800 nM). A cocrystal structure of Hsp90a with 91a (PDB code: 3HEK, resolution: 1.95 Å) established the (R) stereochemistry of the pyrrolidine group as crucial for the molecule to adopt the most favorable, lower-energy conformation to bind at the ATP site. In this configuration, the resorcinol group exhibits a salient H-bond interaction with Asp93, the amide group retains its expected planarity, the 3,3-difluroropyrrolidine group is placed into a hydrophobic region (Tyr137, Val136, Gly135), and the pendant phenyl group is also accommodated in a hydrophobic pocket. Chiral separation of other racemates confirmed the (R)-enantiomers bound far more potently against Hsp90 than their corresponding (S)enantiomers. Unfortunately, all of the (R)-enantiomers exhibited high clearance in human hepatocytes. Armed with this knowledge, a second library was conceived focusing on the synthesis of (R)-enantiomers and the replacement of the phenol groups to improve the ADME profile. The chemistry relied on the enantioselective α -arylation of N-Bocpyrrolidine 92 with (-)-sparteine to prepare chiral 93. Chiral organometallic 93 was coupled with aromatic bromides to afford key intermediate 94. Following Boc deprotection, amines 95 were coupled to a diverse group of aromatic carboxylic acids 96 using HATU to afford library 80. With this methodology enantiomeric ratios of >96:4 were achieved and allowed for the synthesis of a *ca*.112-membered library. Unfortunately, none of the compounds of library 80 exhibited inhibitory activity against Hsp90 when tested at 10 uM in the binding assay. Lastly, selected synthesis of mono-phenol analogs revealed that both phenol groups are essential for binding since the removal of either phenol group led to a >190-fold loss in potency.

Selective Norepinephrine Reuptake Inhibitors.73

The norepinephrine transporter is a membrane bound protein which regulates the uptake of the neurotransmitter norepinephrine (NE) from the presynaptic cleft of noradrenergic neurons during synaptic transmission. It therefore plays an important role in regulating the physiological functions of NE, the deficiency of which has been implicated in a number of neurological disorders. Norepinephrine reuptake inhibitors (NRIs) such as reboxetine and atomoxetine have been used clinically for the treatment of major depressive disorder and attention deficit hyperactivity disorder (ADHD), respectively. Researchers at Pfizer disclosed recently several new chemical series of selective norepinephrine reuptake inhibitors (sNRI).⁷³ The two lead compounds **97** and **98**, investigated clinically, were discontinued due to hepatotoxicity (Figure 6). This safety issue identified with **97** and **98** was thought to be related to the high lipophilicity of these compounds (clogP = 4.2–4.4). A search for new NRI templates, providing ligands with reduced lipophilicity when compared to **97** or **98**, was then undertaken. The benzamide derivative **99** was identified, from

previous lead identification work, as a weak norepinephrine reuptake inhibitor with low selectivity over the serotonin transporter (K_i (NET) = 294 nM; K_i (SERT) = 654 nM). The lead optimization objective was to improve the affinity and the selectivity for the norepinephrine transporter, while simultaneously reducing the lipophilicity in this series. Analogs of **99** (205-member library; general structure **100**) were readily available *via* solution phase parallel synthesis methodology. The N-alkyl or N-aryl substituted 3-amino-1-Boc-(S)-pyrrolidines 102, obtained from 3-amino-1-Boc-(S)-pyrrolidine using common methodologies, were converted to the target compounds 100 via amide formation followed by N-Boc deprotection. The binding affinity of the library compounds at the human norepinephrine, serotonin, and dopamine transporters was determined using scintillation proximity assay (SPA) technology. SAR analysis in this series revealed that potent norepinephrine reuptake inhibition could be achieved over a range of lipophilicity (clogP: 3.0–4.4). The lead optimization campaign led to the discovery of compound **101**, a potent norepinephrine reuptake inhibitor ($K_i = 6 \text{ nM}$) displaying 37-fold and 380-fold selectivity over the serotonin and dopamine transporters, respectively. The lipophilicity of 101 (clogP = 3.3) was also significantly reduced when compared to previous lead compounds 97 and 98 investigated clinically (clogP = 4.4 and 4.2, respectively). Further studies demonstrated that 101 has excellent metabolic stability in human liver microsomes and human hepatocytes, weak CYP (1A2, 2C9, 2C19, 2D6, 3A4) inhibition, and good membrane permeability. In light of its weak inhibitory activities at the hERG (IC₅₀ >20,000 nM) and NaV_{1.5} (IC₅₀ > 26,000 nM) ion channels, 101 is expected to have a satisfactory cardiovascular safety profile. Off-target profiling of **101** at a panel of 110 receptors, enzymes and ion channels revealed only weak affinity at the M₄ ($K_i = 4,300$ nM) and M₅ ($K_i = 1,800$ nM) muscarinic receptors. In vivo rat microdialysis experiments demonstrated that 101 produces a rapid increase in NE levels in the prefrontal cortex after subcutaneous administration demonstrating good CNS penetration. Based on its favorable profile. **101** (PF-3409409) was selected for preclinical evaluation.

Positive Allosteric Modulators of mGluR4.²⁸⁷

Metabotropic glutamate receptors (mGluRs) play important roles in a broad range of central nervous system functions and have therapeutic potential in a variety of neurological and psychiatric disorders. Activation of metabotropic glutamate receptor 4 (mGluR4) has been shown to modulate neurotransmission in the basal ganglia and results in antiparkinsonian effects in rodent models of Parkinson's Disease (PD). Allosteric binding sites, as opposed to traditional orthosteric binding sites, offer unparalleled opportunities for drug discovery by providing high levels of selectivity, mimicking physiological conditions and affording fewer side effects. VU0155041 **103**, a novel mGluR4 positive allosteric modulator (PAM) discovered in a high throughput screen was the starting point of a detailed SAR analysis of this compound class (Figure 7). Three parts of the lead molecule were explored separately through an iterative parallel synthesis approach, the aromatic amide moiety, the carboxylic acid residue and the cyclohexyl core. The first library centered around commercially available *cis*-1,2-cyclohexanedicarboxylic anhydride **104**, which was reacted with respective R¹-amines in THF at 55 °C to give library compounds **105**. Several 3,5- and 3,4-substituted phenyl, unsubstituted phenyl, benzyl and pyridyl, morpholino and cyclo alkyl analogs were

evaluated. Besides the original 3,5-dichlorophenyl lead **103** (EC₅₀ hmGluR4 = 0.74 μ M), only the 3-Cl-5-F-phenyl amide **109** retained some activity (EC₅₀ hmGluR4 = 2 μ M), all other substitutions led to a loss of activity. Carboxylic acid replacements were investigated next, starting from VU0155041 **103**, coupling respective R²,R³-amines with EDC, HOBt and DIEA in DMF a library of diamides **106** was synthesized. All compounds showed at least a tenfold loss of activity, with the exception of the primary carboxamide **112**, which retained similar submicromolar activity as the lead (EC₅₀ hmGluR4 = 0.95 μ M). Variations of the core structure were then examined. Commercially available cyclic anhydrides **107** were reacted with 3,5-dichloroaniline in THF at 55 °C. Only the cyclohexene analog **115** retained some activity (EC₅₀ hmGluR4 = 2.7 μ M), substituted cyclohexene cores (e.g. **117**) resulted in inactive compounds. Changing the substitution pattern to a 1,3-orientation led to a total loss of activity. Combining the cyclohexene core with a primary carboxamide in compound **116** showed some activity (EC₅₀ hmGluR4 = 3.1 μ M).

This SAR evaluation around VU0155041 **103**, utilizing a parallel synthesis approach, illustrates another example of the rather flat SAR common to positive allosteric modulators.

mGluR5 allosteric modulators.⁷¹

Excess dopamine transmission in the brain is believed to be one cause of schizophrenia and antipsychotic agents that antagonize the dopamine D2 receptor are routinely prescribed. Many of these agents display off target pharmacology against a range of neurotransmitters leading to side effects. It was observed clinically that administering glycine, an N-methyl-Daspartate (NMDA) receptor co-agonist, elicited a modest improvement in schizophrenic patients suggesting that activation of the NMDA receptor could be an option for therapeutic treatment. Potentiation of NMDA receptor transmission can occur directly by modulating NMDA receptor sites or indirectly by activating NMDA-receptor-function dependent GPCRs. Metabotropic glutamate receptor 5, mGluR5, is a GPCR that upon activation can potentiate indirectly the NMDA receptor. Using ADX-47273 (118), a previously reported positive allosteric modulator of mGluR5 demonstrating in vivo activity in cognition and schizophrenia animal models, Engers and colleagues generated solution-phase libraries around scaffold 119 to flesh out the SAR of 118 and improve its overall physicochemical properties (Figure 8).⁷¹ Two routes were developed for library synthesis. The first route involved simultaneous coupling-cyclization of (Z)-N'-hydroxyimidamides 120 and (S)-1-(tert-butoxycarbonyl)piperidine-3-carboxylic acid 121 using EDCI and HOBt under reflux to give oxadiazoles. Removal of the Boc group and acylation of the piperidine intermediates afforded library **119a**. The second route consisted of the esterification of (S)-piperidine-3carboxylic acid 122, acylating ester 123, saponifying N-acyl esters 124 with LiOH and then simultaneous coupling-cyclization with (Z)-N'-hydroxyimidamides 120 to afford desired library 119b.

In the first exploratory library, the 4-fluorobenzylpiperamide was kept constant and the 4-fluorophenyl ring on the oxadiazole unit was replaced. It was found that 2-pyridyl and 2-thienyl groups were reasonable substitutes for the 4-fluorophenyl group while the 3- and 4-pyridyl and 2-pyrazinyl groups were not. In the follow-up library **119b**, which retained the original 4-fluorophenyl and now included the 2-thienyl and 2-pyridyl (R¹) groups, the acyl

group (\mathbb{R}^2) was varied. Biological evaluation of this library revealed the following trends. For \mathbb{R}^2 , most mono- and di-fluorophenyl groups exhibited submicromolar ago-potentiation (potentiation of the agonist activity of glutamate at its EC₉₀) while the 2,6-difluorophenyl group did not. The 2-pyridyl analogs e.g. **125**, were 2-times less potent than their 4fluorophenyl and 2-thienyl counterparts. Unexpectedly pyridyl compounds, e.g. **128**, displayed pure mGluR5 positive allosteric modulation as these compounds had no inherent agonist activity. The corresponding HCl salts of the pyridyl analogs exhibited expected improved water solubility. Interestingly, the cyclobutyl analog **129** was a negative allosteric potentiator, an example of an unusual functional switch. Finally, in order to establish the importance of the chiral center, enantiomeric separations were carried out on **118**, **125** and **126**. The (*R*)-enantiomers were in general 9- to 10-fold less potent than the (*S*)-enantiomers, yet display similar in vitro efficacy. With this study the SAR features of **118** were quickly identified, particularly the relevance of the (*S*)-chiral center.

Bcl-2 inhibitors: Diversity-Oriented Synthesis Library.¹⁸²

Apoptosis, or programmed cell death, is important for normal development, host defense, and suppression of oncogenesis, and disregulation of apoptosis has been implicated in cancer and many other human diseases. The Bcl-2 family proteins are central regulators of apoptosis and comprise anti-apoptotic proteins such as Bcl-2, Bcl-xL, and Mcl-1 and proapoptotic proteins such as Bak, Bax, Bim, Bid, and Bad. Overexpression of the Bcl-2 membrane protein has been observed in 70% of breast cancer, 30-60% of prostate cancer, 80% of B-cell lymphomas, 90% of colorectal adenocarcinomas, and many other forms of cancer. The expression levels of Bcl-2 proteins also correlate with resistance to a wide spectrum of chemotherapeutic drugs and y-radiation therapy. Bcl-2 is therefore a promising molecular target for the design of an entirely new class of anticancer drugs aimed at overcoming resistance of cancer cells to apoptosis. Consequently, design of non-peptide small molecule inhibitors of Bcl-2 and Bcl-xL is currently an exciting research area for the development of new anticancer agents. In an effort to identify novel inhibitors of Bcl-2, scientists at Infinity Pharmaceuticals designed and prepared a 15,000-member pyridone library (Figure 9a/b), *via* discovery oriented synthesis (DOS).¹⁸² The design of this library was based on the structure of the nicotinic agonist (-)-cytisine (130) and previous work describing the total synthesis of this natural product. In particular, the pyridone cyclization step, key transformation for the total synthesis of (-)-cytisine, was exploited for the diversity oriented synthesis of heterocycles 131a and 132a as well as their corresponding enantiomers 131b and 132b, respectively. The synthons 140 and 141 used for the diversity oriented library synthesis were prepared according to Figure 9a. Condensation of D-glyceraldehyde acetonide 133 with phosphonate 142, under Horner-Emmons conditions, provided the α . β unsaturated ester 134, which reacted with the imine 135 in toluene in the presence of AgOAc and DBU to provide the corresponding [3+2] azomethine ylide-alkene cycloaddition product 136 in high yield and excellent diastereoselectivity (>95:5). The pyrrolidine derivative **136** was then converted to the bridge bicyclic scaffold **140** in 5 steps, i.e. a) protection of the NH functionality of 136 as the N-Fmoc derivative, b) deprotection of the allyl ester functionality by treatment with $Pd(PPh_3)_4$, c) conversion of the resulting carboxylic acid to the corresponding primary alcohol, d) activation of the resulting alcohol

with mesyl chloride followed by formation of the pyridone ring via intramolecular cyclization, and e) conversion of the 1,3-dioxolane, 2,2-dimethyl functionality to the corresponding primary alcohol according to a 3-step sequence. The tricyclic synthon 141 was prepared from phosphonate 143 via a synthetic sequence similar to the one described for the preparation of 140. The pyridone scaffolds 140, 141, prepared in large scale (>75g), were then loaded onto silicon-functionalized Lanterns via activation with TfOH (average loading level of 15 mmol of compound per Lantern). Deprotection of the N-Fmoc functionality provided the resins 144 and 148 which were converted to the library compounds 145-147 and 149-151 via classical solid-phase derivatization methodologies (Figure 9b). The 15,000 library compounds, prepared with purities greater than 75% for over 75% of the library, were screened for binding affinity for Bcl-2 and Bcl-xL. The most potent compounds derived from the bridged bicyclic pyridone scaffolds 131a and 131b, exemplified by compound 152 [K_i (Bcl-2) = 2.0 μ M; K_i (Bcl-xL) = 5.7 μ M)] and its enantiomeric analog 153 [K_i (Bcl-2) = 1.3 μ M; K_i (Bcl-xL) = 6.6 μ M)], contained a chlorosubstituted diphenyl 2-aminothiazole and a diamine at the R and R' position, respectively. Compound 154 was the best ligand identified from the tricyclic pyridone cores 132a and **132b.** This compound selectivitly binds with micromolar affinity at Bcl-2 ($K_i = 1.2 \mu M$) and displays, in contrast to 152,153, high (> 100-fold) selectivity for Bcl-2 over Bcl-xL (Figure 9b).

Catch and Release Synthesis of Substituted Guanidines.²⁸⁹

One of the most common methods to prepare substituted guanidines 156 involves the condensation of amines with S-methylated thioureas 156 (Figure 10). This method suffers from the formation of noxious and toxic methyl mercaptan side product, which limits application for high-throughput parallel synthesis. To overcome this problem, scientists at Abbott designed a solid phase strategy for the synthesis of N, N', N''-substituted guanidines using a catch and release methodology inspired from the solution phase synthesis described above.²⁸⁹ In this method, the thioureas 155 immobilized on solid support react with various amines to provide the desired guanidine derivatives, free of mercaptan side products. Using this new methodology, and after several optimization of the reaction conditions, various substituted guanidines were prepared from thioamides in a one-pot process. Hence, loading of N-substituted thioamides 157 to brominated polystyrene resin in a mixture DCM/DMF (2:1) at 50°C for 4–6 h provided the thiourea resins 155, which could be washed, dried, and stored or used without isolation for the next step. Condensation of the resin-bound thioureas with primary amines at 50 °C for 20 h provided, in high yield, the desired substituted guanidines 156a,b purified by HPLC and isolated as their TFA salts. The reaction conditions were modified to allow the formation of N, N', N''-trisubstituted guanidines from the corresponding secondary amines in good yields. In the modified method, the condensation of secondary amines to the resin-bound thioureas was conducted in methoxyethanol in the presence of 1.5 equiv. of HgCl₂. Using this new solid phase methodology, a total of $17 N_{,N'}$ disustituted or N, N', N''-trisubstituted guanidines **156a,b** were prepared, from the corresponding thioamides, in isolated yields ranging from 42% to 99%.

Substituted Pyrimidines via a 3-component reaction.¹⁵⁸

Konakahara and coworkers developed a novel three-component coupling reaction of enamines 158, triethyl orthoformate and ammonium acetate providing a facile route to 4,5disubstituted pyrimidine derivatives 160 (Figure 11). Pyrimidine derivatives exhibit biological activity against all major classes of molecular targets including kinases, proteases, nuclear hormone receptors and GPCRs. Bredereck- and Pinner-type chemistries are established routes for the preparation of pyrimidines. However, challenging multistep synthesis of reactive intermediates and harsh reaction conditions limit their general utility, particularly for the preparation of pyrimidine-based arrays. The initial 3-CR was conducted by simply heating enamines 158 with 3 equiv of orthoester and 2 equiv of NH₄OAc in toluene at 100 °C for 20 h. Optimized reaction conditions included the addition of 0.1 equiv of ZnCl₂ which improved the overall yield (>70%), accelerated reaction time and enhanced the range of substrates. Non-commercially available enamines were prepared in high yield (>90%) by the addition of electron-withdrawing stabilized carbon nucleophiles to aryInitriles. The ZnCl₂-catalyzed three-component reaction was expanded by using ketones 159 in place of enamines. The 3-CR with ketone substrates required 72 h for completion affording the desired products in good yields (54-70%).

Discovery of Ned-19.340

Second messengers play enormously important roles in transduction of cellular signals and govern cellular responses to stimuli. Molecular and cellular biology has benefited greatly from molecular tools like Forskolin which alters cellular levels of the critical second messenger cAMP. Nicotinic acid adenine dinucleotide phosphate (NAADP, 161) is recognized as a critical regulator of Ca²⁺ release in numerous human tissues in response to several reported stimuli (Figure 12). Controversies surrounding the exact nature of NAADP as a secondary messenger and its mechanism of action permeate the literature. Furthermore, the lack of chemical probes of this important biomolecule hamper studies into the cause and consequences associated with Ca²⁺ release. To rectify this inadequacy, Churchill and coworkers reported the discovery of Ned-19 (162), a potent inhibitor of NAADP signaling.³⁴⁰ Utilizing the chemical structure (including electrostatic surface assessments) of NAADP, a ligand-based virtual screen was performed in which novel structures were sought with sufficient overlap with NAADP while maintaining drug-like properties in the 'hit' compounds. One agent that was discovered was an (S)-2,3,4,9-tetrahydro-1H-pyrido[3,4blindole-3-carboxylate designated as Ned-19. This commercially available agent inhibited NAADP signaling in a sea urchin egg assay at nanomolar concentrations. The stereochemistry at the 1 position of the tetrahydro-1H-pyrido ring was explored via an independent synthesis. The synthesis was initiated by installment of a chloromethyl group at the 3-position of commercially available 4-methoxybenzaldehyde. The resulting aldehyde was subjected into a Pictet-Spengler reaction with (S)-methyl 2-amino-3-(1H-indol-3yl)propanoate to afford the core (S)-2,3,4,9-tetrahydro-1H-pyrido[3,4-b]indole-3carboxylate scaffold. Displacement of the benzyl chloride with 1-(2-fluorophenyl)piperazine and saponification of the methyl ester provided Ned-19. The diastereomeric mixture was separated following the Pictet-Spengler reaction. The specific activity of both the *cis* and trans isomers of Ned-19 were evaluated and the trans isomer was significantly more active

than the *cis* isomer. Biochemically, Ned-19 was able to eliminate all NAADP-mediated Ca^{2+} signaling at 100 µM without altering inositol 1,4,5-trisphosphate-mediated or cADP-ribose-mediated Ca^{2+} release. These experiments suggest that Ned-19 is selective as a NAADP signaling probe rather than a nonspecific modulator of Ca^{2+} release. Ned-19 blocks NAADP signaling in intact cells and it can serve as a fluorescent label for NAADP receptors. Finally, Ned-19 was used to demonstrate a role for NAADP signaling in glucose signaling and Ca^{2+} release in pancreatic beta cells. While further studies with Ned-19 are required to fully validate its use as a probe of NAADP signaling, this work offers a novel agent to begin to fully dissect the various roles of this important secondary messenger.

Discovery of DG-041.338

There exists a complex network of signaling events that govern platelet activation and aggregation and great strides have been made in modulating these relevant signaling events to avoid catastrophic acute thrombosis during vascular events like myocardial infarction and stroke. The recognition that aspirin (acetylsalicylic acid) acts as an antagonist of platelet aggregation can be considered a starting point for research into defining the critical targets associated with platelet function. More recently, the purinergic P2Y12 receptor has been shown to play an essential role in platelet signaling following platelet collagen and von Willebrand factor release and Ca²⁺ stimulation of TxA₂ and release of ADP. P2Y₁₂ receptor antagonists including prasugrel have proven more effective than aspirin in preventing recurrent myocardial infarction. Both aspirin and $P2Y_{12}$ receptor antagonists, however, have the unwanted side-effect of severe, prolonged bleeding events due to their global impairment of platelet aggregation. Research has continued to shed light on the complex signaling cascade associated with platelet aggregation and today it is known that multiple GPCRs are responsible for both Ca²⁺ release and activation/inactivation of adenylcyclase. One target that is gaining attention is the EP₃ receptor. The EP₃ receptor responds to prostaglandin E₂ (PGE₂) which is produced in low concentrations within atherosclerotic plaque. Importantly, signaling through the EP₃ receptor is not sufficient to produce an aggregation response without co-signaling events at one or more additional receptors that are associated with platelet activation. Studies have showed that platelets lacking the EP₃ receptor were protected from thrombotic events in several models and it has been hypothesized that, since healthy tissues do not produce PGE₂, blocking of the EP₃ receptor should have a diminished effect on bleeding events. To examine this theory, researchers at deCODE chemistry sought out new EP₃ receptor antagonists through a ligand-based design strategy (Figure 13).³³⁸ The basic strategy was to seek out scaffolds that mimic the three-dimensional shape and electrostatic profile of the native ligand PGE₂. Ultimately efforts focused on a collection of substituted indoles with appendages at the 1 and 7 positions. Their initial leads possessed low-nanomolar activity at the EP₃ receptor and lead optimization included blocking metabolically labile positions at C3 and C5 of the indole ring. The optimization campaign led to the discovery of DG-041 (163). DG-041 (163) displayed IC_{50} values of 8.1 nM and 4.6 nM in a Ca²⁺ response assay and a radioligand displacement assay, respectively. DG-041 had no relevant activity versus the functionally related, platelet-associated GPCR targets nor was this agent active across a panel of 50 random GPCRs. DG-041 was also found to be acceptable for both oral and i.v. dosing with appropriate PK properties for use as

a tool compound in models of platelet aggregation. The synthesis of this agent was accomplished via sequential intramolecular and intermolecular Heck couplings to arrive at an appropriately 7-substituted indole intermediate. Saponification followed by coupling to 4,5-dichlorothiophene-2-sulfonamide and subsequent *N*-alkylation with 2,4-dichloro-1- (chloromethyl)benzene provided DG-041. DG-041 was found to inhibit human and rat platelet aggregation in vitro in a collagen induction experiment with several co-agonists and, importantly, in the presence of high serum concentrations. Within *in vivo* systems, DG-041 inhibited platelet aggregation at a dose of 10 mg/kg while there was no prolongation of bleeding up to a concentration of 100 mg/kg. The authors present several other findings that further validate DG-041 as an important new tool for studying the EP₃ receptor and reference, as yet, unpublished data suggesting that DG-041 is efficacious in a human patient population.

Displaceable fluorous dihydropyran linker.322

Fluorous linker combined with fluorous solid-phase extraction (F-SPE) is a powerful reaction and purification technique for solution-phase synthesis.³⁸⁰ The Nelson group introduced fluorous dihydropyran linkers for the protection of amino and hydroxyl groups. Figure 14 highlights the utility of displaceable linker **164** in the synthesis of highly condensed heterocyclic compound **165**.³²² The fluorous linker was attached to sulfonamide **166** through Fukuyama-Mitsunobu reaction to form compound **167**. Cascade metathesis using Hoveyda-Grubbs second generation catalyst cleaved the central dihydropyran ring and produced compound **168**. This compound was then used for Diels-Alder reaction with 4-phenyl-[*1,2,4*]-triazole-3,5-dione **169** to afford **170** as a single diastereomer. The removal of the *o*-nitrophenylsulfonyl (Ns) group from **170** followed by the reaction with an isocyanate gave urea **171**. Finally, the fluorous linker was removed by the treatment of 3% TFA to afford product **165**. In this multistep synthesis, all the intermediates and the final product were isolated from the reaction mixture by F-SPE.

Displaceable fluorous isonitrile linker.³²⁷

Isonitriles are versatile reagents for organic reactions. However, their synthetic advantages are offset by the notorious odor associated with conventional isonitriles. The Pirrung group developed a new process using base-promoted ring-opening of oxazoles to produce aromatic isonitriles which no longer have the unpleasant odors.²²⁸ During the reaction process a fluorous sulfonate group could be introduced to form a fluorous linker. The utility of the displaceable fluorous isonitrile has been demonstrated in the multistep synthesis involving Ugi four-component reaction (U-4CR) and metal catalyst-promoted linker cleavage reactions. Freshly prepared isonitrile **172** was used as the limiting agent for the Ugi reaction to form **173** (Figure 15). This intermediate was then subject to different coupling reactions to remove the fluorous linker and also to introduce a new functional group to the Ugi condensation product. The developed post-Ugi reactions included Suzuki coupling to form compound **176**. The protocol is suitable for diversity-oriented synthesis of compound libraries since four substitution groups can be introduced during the Ugi reactions and different linker cleavage reactions could lead to an array of scaffolds.

Fluorous synthesis of 1,4-benzodiazepine-2,5-dione library.³²⁷

In the solution-phase parallel synthesis of a 1,4-benzodiazepine-2,5-dione library, the Yang group developed a new U-4CR using fluorous benzaldehydes instead of fluorous isonitriles (Figure 16) as the displaceable linker.³²⁷ The library synthesis was accomplished in three reaction steps. At the first step of the Ugi reactions, fluorous benzaldehydes **177** were used as the limiting agent to react with Boc-anthranilic acids **178**, amino esters **179**, and cyclohexyl isonitrile **180**. After F-SPE purification, Ugi products **181** were used for the second step of acetyl chloride-promoted de-Boc/cyclizations to form 1,4-benzodiazepine-2,5-diones **182**. The cyclization reactions were selective and only attacked the ester group to form the seven-membered ring. The last step Suzuki reactions were carried out under microwave heating to cleave the fluorous sulfonyl group and introduce the biaryl functional group to form products **183**. The Yang group also used 2-nitrobenzoic acids as anthranilic acid alternatives for the U-4CRs. A demonstration library of 36 analogs was produced with building blocks R¹ to R⁴ variants.

Fluorous synthesis of piperazinedione-fused tricyclic compound library.³²⁸

Fluorous amino esters are useful linkers for library synthesis. In a joint work conducted by the Werner and the Zhang groups, fluorous amino esters **184** were used for solution-phase parallel synthesis of novel piperazinedione-fused tricyclic library compounds **189** (Figure 17).³²⁸ The compounds are structurally similar to the tricyclic thrombin inhibitors and diketopiperazine-based inhibitors of human hormone-sensitive lipase. At the first library synthesis step, azomethine ylides generated from **184** and aldehydes **185** underwent microwave-assisted 1,3-dipolar cycloadditions with maleimides to form **186** in stereoselective fashion. The adducts were treated with chloroacetyl chloride to afford *N*-acylated products **187**. The chloro group of **187** was displaced with amines to afford **188**. The last step was the microwave-promoted cyclizations to cleave the fluorous linker and generate the piperazinedione ring. The final products were purified by HPLC to ensure >95% purities. Library **189** containing 90 compounds was synthesized with assorted building blocks for R¹ to R⁴.

Fluorous mixture synthesis of natural product resorcylic acid lactone (RAL) library.³²⁹

Natural resorcylic acid lactones (RAL) containing a *cis*-enone moiety such as radicicol A, L-783277, and LL-Z1640-2 are known to be potent and irreversible kinase inhibitors. For QSAR studies, the Winssinger group developed a fluorous mixture synthesis (FMS) method based on the previously reported fluorous total synthesis of radicicol A and synthesized a compound library containing 51 analogs (Figure 18).³²⁹ In the FMS, three propargyl alcohols were used each attached to a different *p*-methoxybenzyl (PMB) linker by reacting with F-PMB-trichloroacetimidates **190** to form **191(a–c)** (Scheme 5). The equimolar mixture of **191(a–c)** was split to three portions and each reacted with one of three aldehydes **192(a–c)** to afford 3 mixtures of **193(a–c)**. After sequential hydroxyl group protection, TBDPS deprotection, and iodination, compounds **193(a–c)** were converted to 3 mixtures of **194(a–c)** each containing three components. Through a similar pathway, compound **191d** reacted with

aldehydes 192(a-c) to afford compounds 194(d-f). The second stage of FMS involved the reaction of three compounds of 195(a-c) with each of 6 compounds of 194(a-f) to afford compounds 196. Among the 18 pools of 196, 9 are three-component mixtures generated from **194(a–c)** and 9 are single compounds generated from **194(d–f)**. At this point, those 9 three-component mixtures were demixed on HPLC with a fluorous column to afford 27 individual compounds of 196. These 27 compound and another 9 compounds of 196 generated from 194(e-f) were treated with DDQ and TBAF to remove both the fluorous linker and the TMSE group. The macrocyclization reactions were promoted by Mitsunobu reaction conditions using fluorous PPh₃ and DIAD. The fluorous derivatives were easily removed from the reaction mixture by F-SPE. The treatment of NaOH removed the Bz group to afford total of 36 macrocyclic compounds **197**. Among them, $24 (X = CH \text{ or } CH_2)$ were treated with DMP followed by HF to afford products 199. Another 12 of 197 were treated with PS-SO₃H and PS-IBX to afford products 200. Compounds 199 and 200 can be converted to 201-205 through simple transformations to afford a total of 51 macrocyclic compounds. A subset of 28 representative library compounds was assayed against a panel of 19 kinases representing those bearing the adequately positioned cysteine residue, bearing a cysteine residue at a different position within the ATP binding pocket, and those that do not bear a cysteine residue. The screening results indicated that there is very little difference in relative selectivity of kinase inhibition throughout the library compounds. VEGF-R2 is the most highly inhibited kinase followed by PDGFR-a, VEGR-R3, Flt3, VEGF-R1, MEK1 SESE and KIT. Two representative compounds 199a and 200a were evaluated in a larger panel of 402 kinases and also evaluated against a series of mutations of Fli3 and KIT. The screening results provided valuable QSAR information and also led to the identification of several potent inhibitors of multiple oncogenic kinases.

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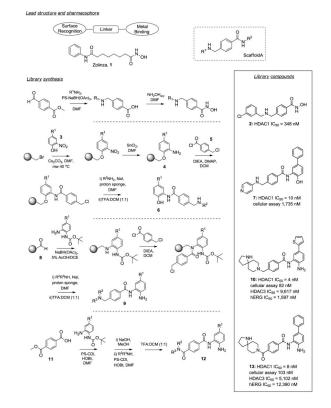


Figure 1. HDAC1/HDAC2 inhibitors.¹³⁴

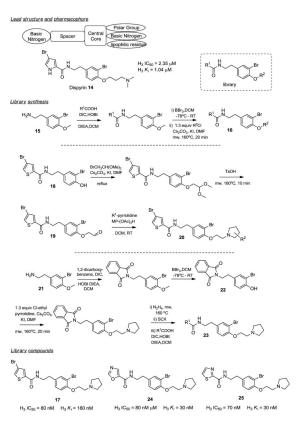


Figure 2. Histamine H₃ antagonists.¹³⁶

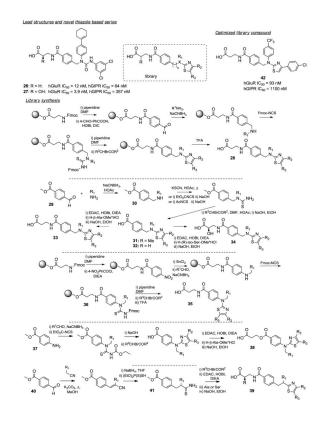


Figure 3. Glucogon receptor antagonists.¹⁷⁹

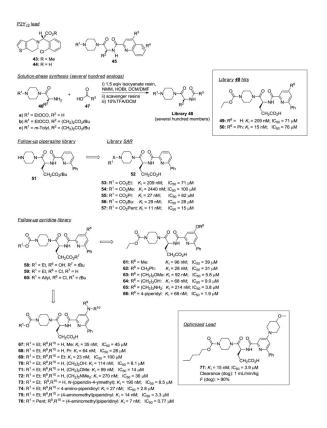


Figure 4. P2Y₁₂ receptor antagonists.²²¹

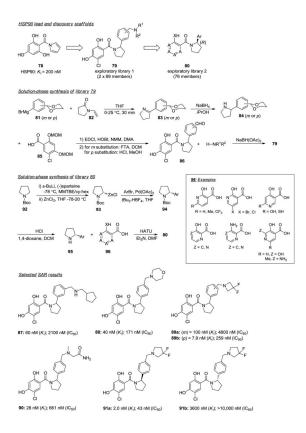


Figure 5. Hsp90 inhibitors.⁴¹

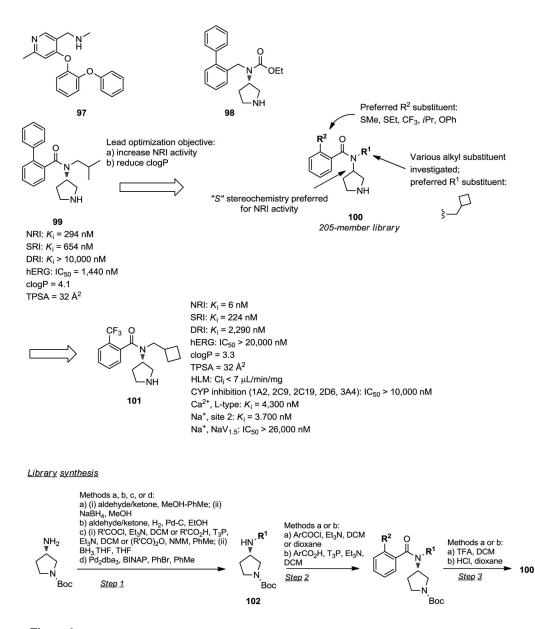


Figure 6. Selective noradrenaline reuptake inhibitors.⁷³

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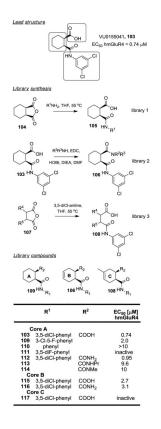
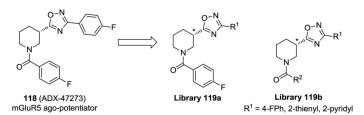
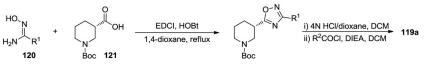


Figure 7. Positive allosteric modulators of mGluR4.²⁹⁷

Lead & SAR scaffold strategy



Solution-phase library synthesis: route 1



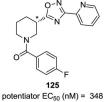
Solution-phase library synthesis: route 2



SAR results:

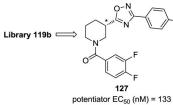
 EC_{50} values determined at an EC_{20} concentration of glutamate Glu Max (%) is the maximum response of compounds relative to the maximal glutamate response

Library 119a 🗆 118 potentiator EC₅₀ (nM) = 168





potentiator EC₅₀ (nM) = 170 Glu Max (%) = 98



Glu Max (%) = 105

Glu Max (%) = 107



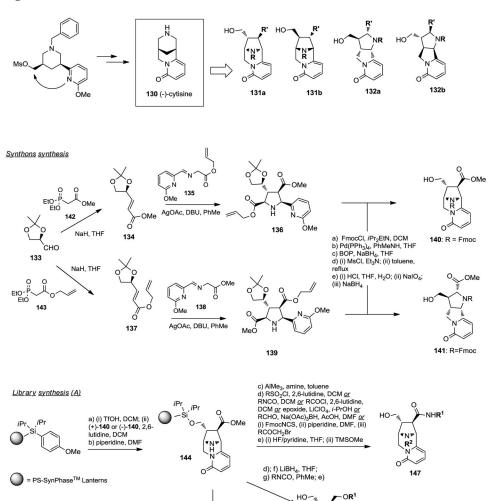
Glu Max (%) = 109

potentiator EC_{50} (nM) = 244 Glu Max (%) = 106 Glu Max (%) = 23%

129 potentiator EC_{50} (nM) = 8700

Figure 8. mGluR5 allosteric modulators.71

Figure 9a.



d); h) LiOH, THF, MeOH, H₂O; e)

OH

N

145

HO

R² =

2-H 2-S=0

146

0-

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Figure 9b.

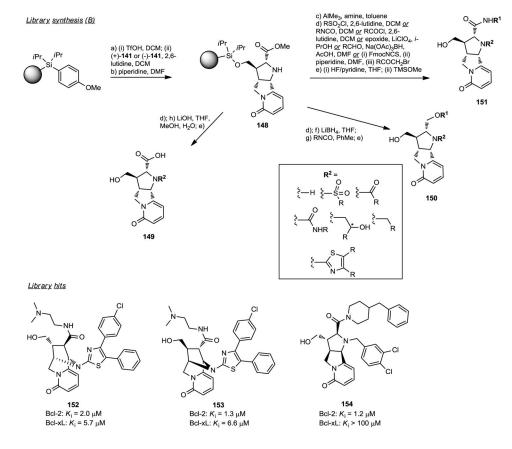


Figure 9.
Figure 9a. Bcl-2 inhibitors via DOS library.¹⁸²
Figure 9b. Bcl-2 inhibitors via DOS library (continued).

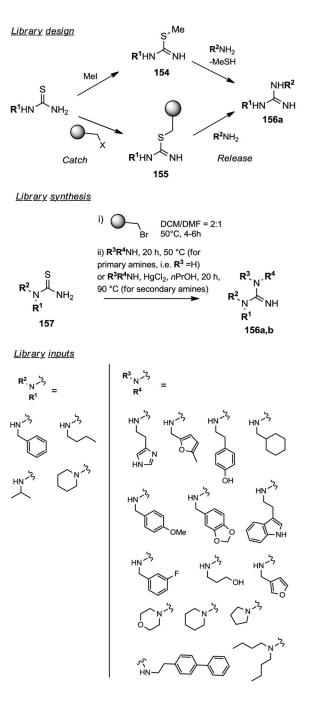
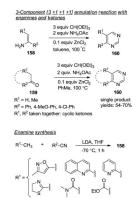


Figure 10. Catch and release synthesis of substituted guanidines.²⁸⁹



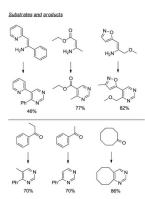
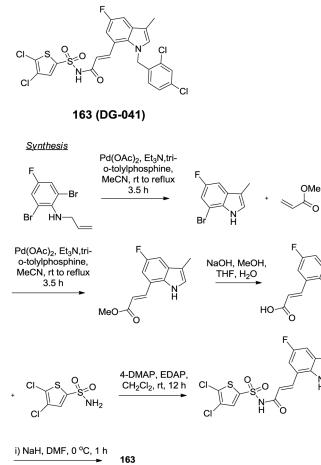
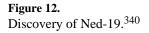


Figure 11. MCR to pyrimidine derivatives.¹⁵⁸

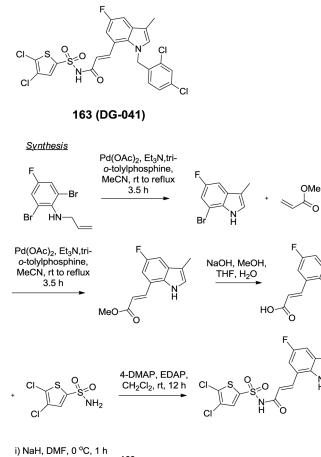
ò



ii) 2,4-dichloro-1-(chloromethyl)benzene, rt, 12 h



ò



163 ii) 2,4-dichloro-1-(chloromethyl)benzene, rt, 12 h

Figure 13. Discovery of DG-041.338

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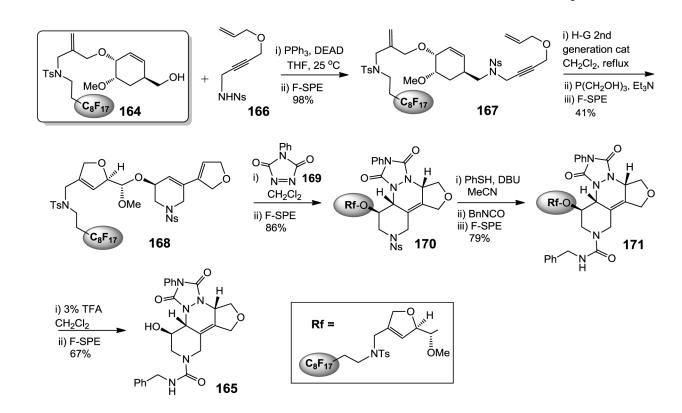


Figure 14. Fluorous dihydropyran linker.³²²

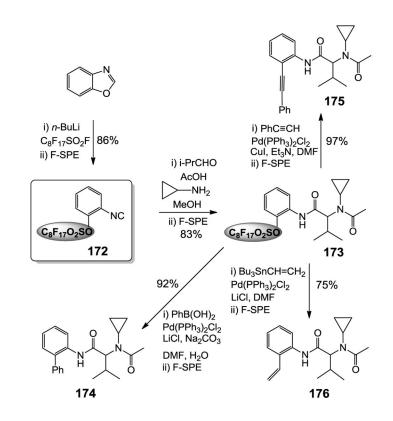
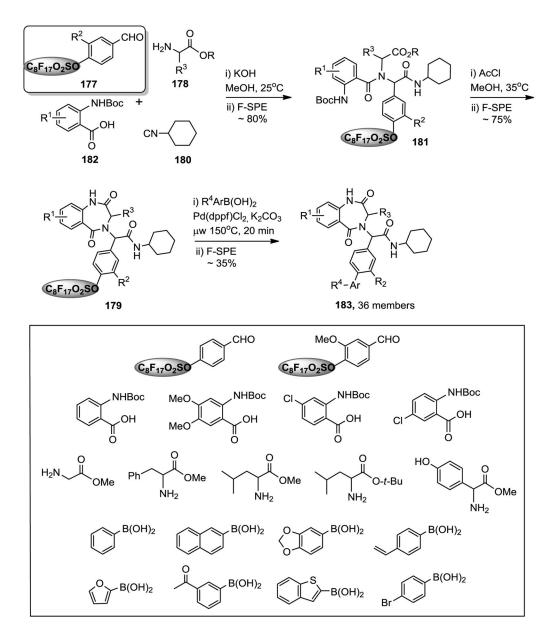


Figure 15. Fluorous isonitrile linker.³²⁷





Fluorous synthesis of 1,4-benzodiazepinedione library.³²⁷

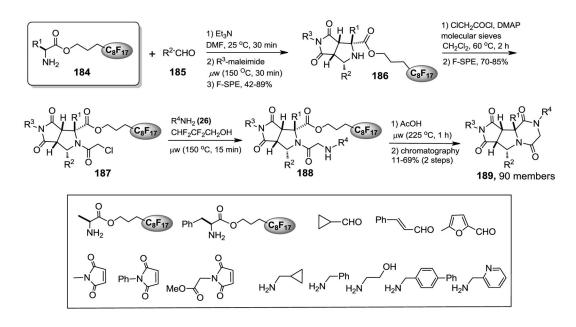
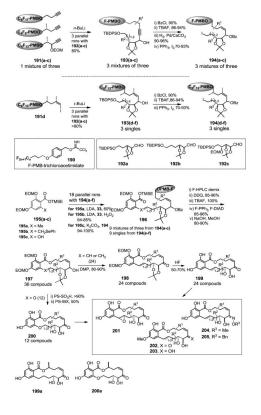
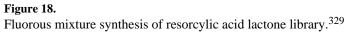
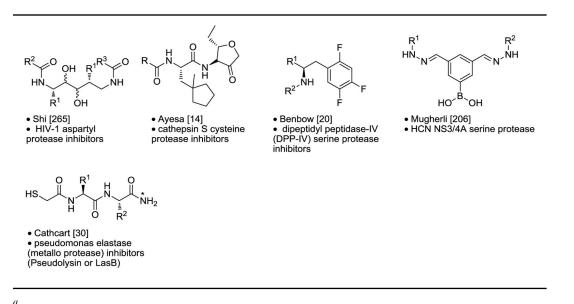


Figure 17. Fluorous synthesis of tricyclic library.³²⁸



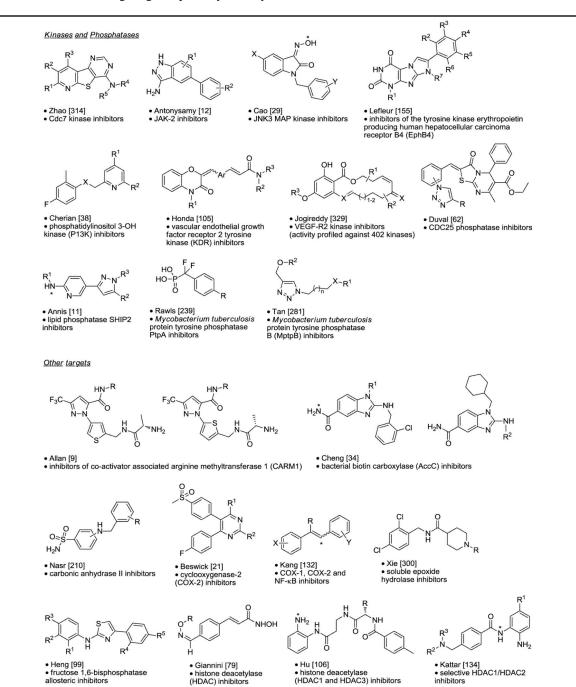


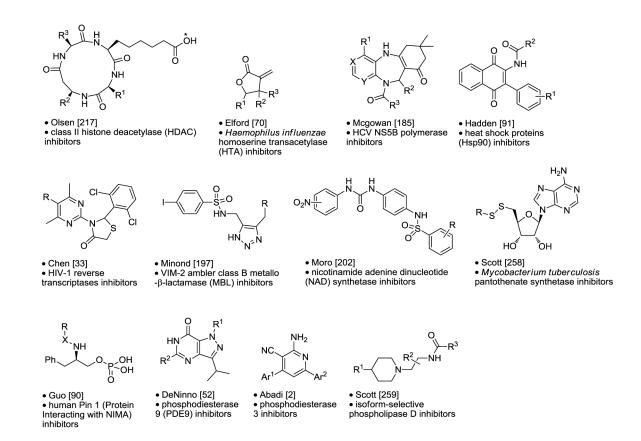
Chemical Libraries Targeting Proteases.



^{*a*}Asterisk is the point of attachment to resin.

Chemical Libraries Targeting Nonproteolytic Enzymes.

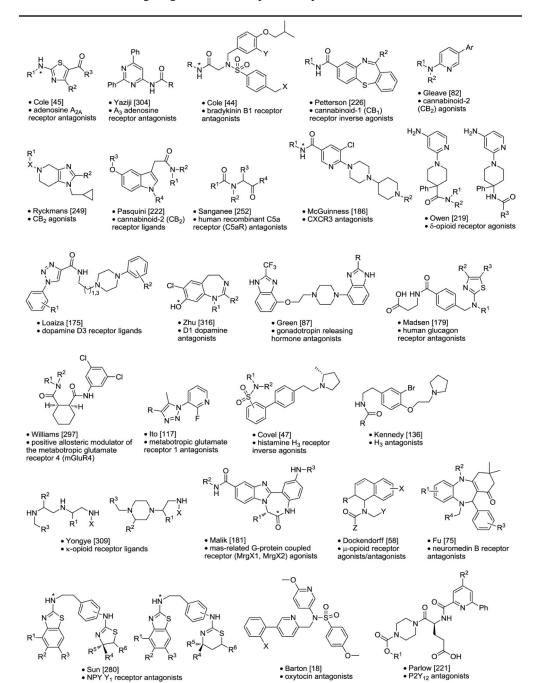


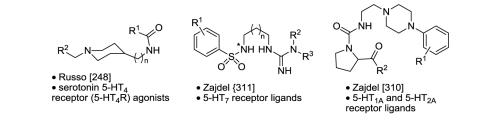


 a Asterisk is the point of attachment to resin.

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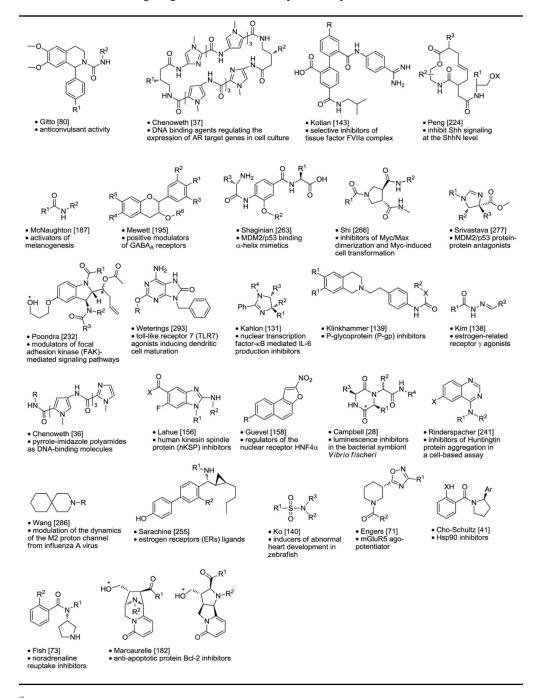
Chemical Libraries Targeting G-Protein Coupled Receptors.





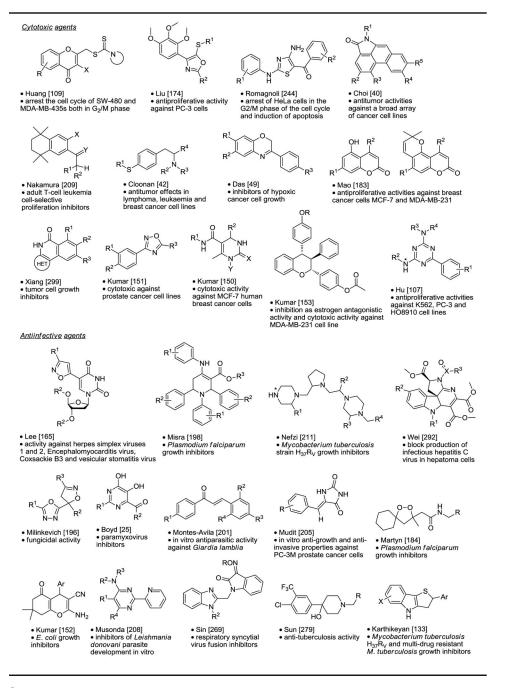
^aAsterisk is the point of attachment to resin.

Chemical Libraries Targeting Non-G-Protein-Coupled Receptors.



^aAsterisk is the point of attachment to resin.

Chemical Libraries Yielding Cytotoxic and Antiinfective Agents.



^aAsterisk is the point of attachment to resin.

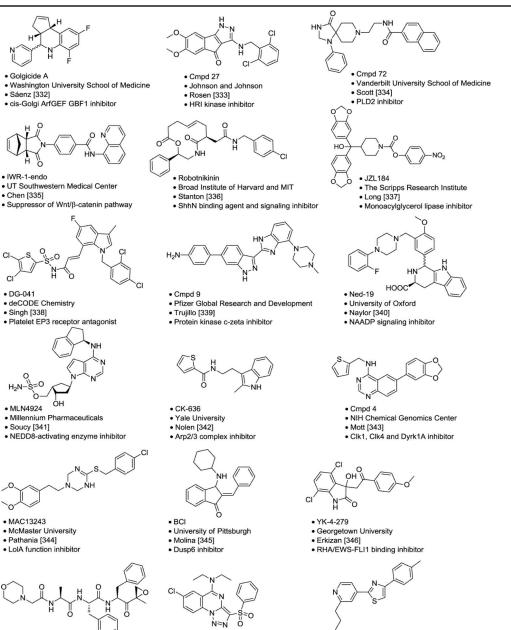
CI

H₂N

0

Table 6

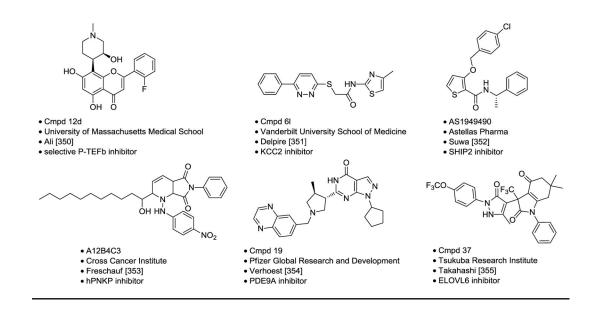
Selected Molecular Probes.



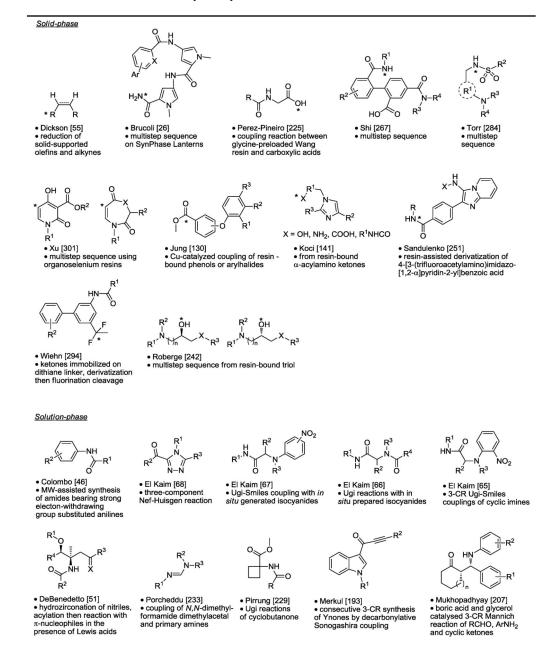
• PR957 Proteoliz, Inc.

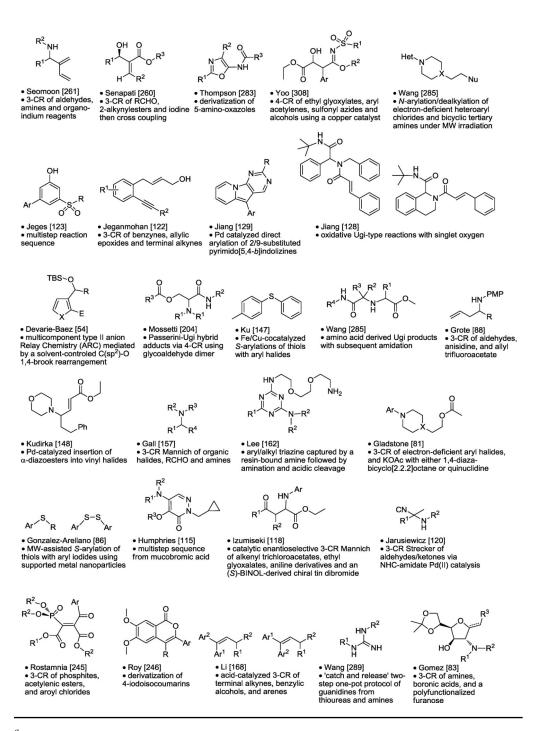
- Muchamuel [347] LMP7 inhibitor
- 1835F03
- Harvard Medical School Swoboda [348]
- Teichoic acid biosynthesis inhibitor
- Kyoto University and Baylor College of Medicine
- Kamisuki [349]
 inhibitor of SREBP activation

Fatostatin



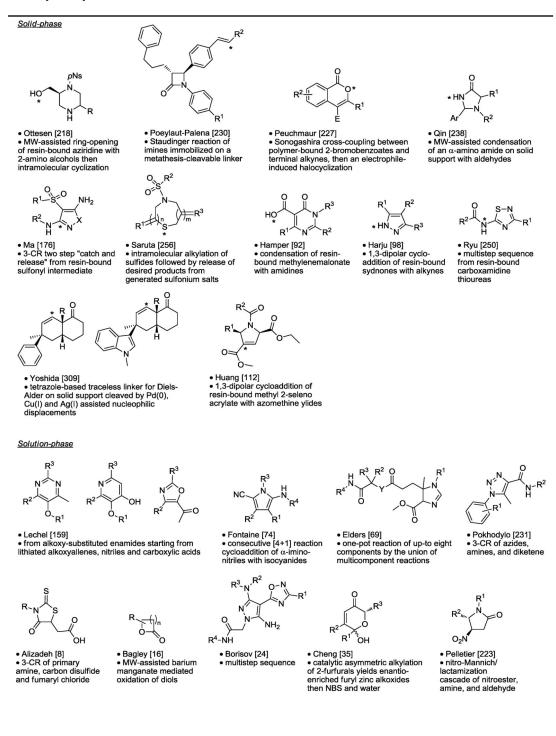
Scaffold Derivatization and Acyclic Synthesis.

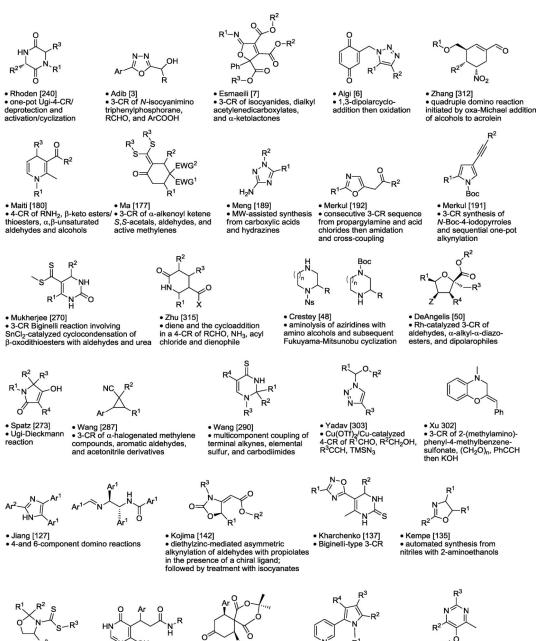




^aAsterisk is the point of attachment to resin.

Monocyclic Synthesis.





- Han [93]
 4-CR of aminoethanols, ketones, carbon disulfide, and halides
- - Hao [96]
 MW-assisted reaction of arylidene-Meldrum's acid, 6-hydroxypyrimidin-4(3H)-one, and amines

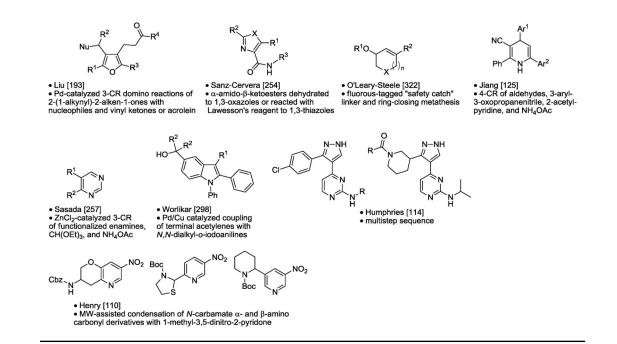
J Comb Chem. Author manuscript; available in PMC 2014 August 21.

Harju [97]
multistep sequence from *N*-acylated amino acids



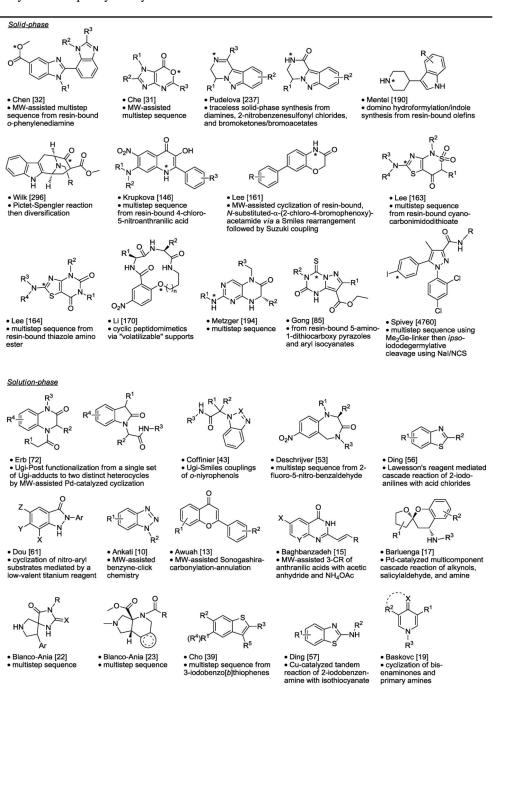
Lechel [160]
3-CR of lithiated alkoxyallenes, nitriles, and carboxylic acids then condensation with ammonium salts

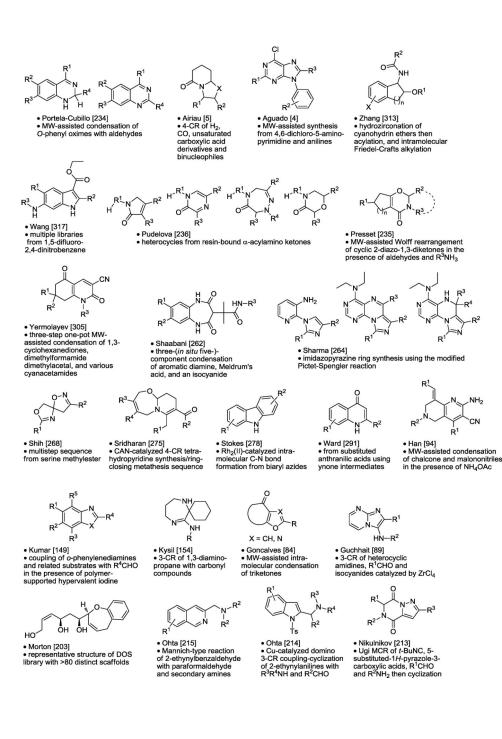
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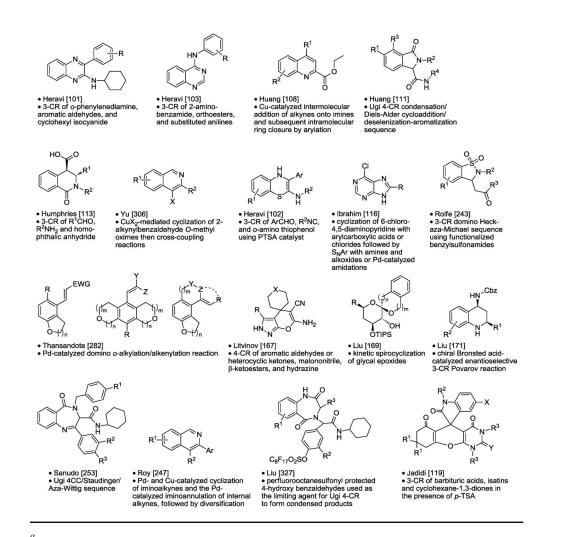


^aAsterisk is the point of attachment to resin.

Bicyclic and Spirocyclic Synthesis.







^aAsterisk is the point of attachment to resin.

Solid-phase

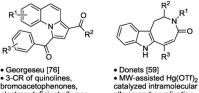
Table 10

Polycyclic and Macrocyclic Synthesis.

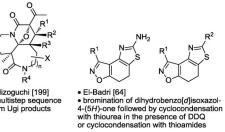


 Huang [110]
 condensation of resin-bound
 3-(2-aminophenylamino)-2 seleno-ester with isothiocyanates and a-amino-acids

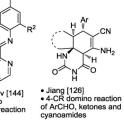
Solution-phase







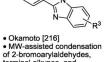
 Mizoguchi [199] multistep sequence from Ugi products



• Kouznetsov [144] • 3-CR imino Diels-Alder reaction



 Hermange [104]
 diastereoselective 3-CR
vinylogous Mannich between isoquinolines, acyl/sulfonyl chlorides, and silyloxyfurans



R

NH₂

alkyne carbocyclization

of 2-bromoarylaldehydes, terminal alkynes, and 1,2-phenylenediamines

R R

HN

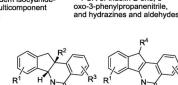
• two tandem isocyanide-based multicomponent

Krasavin [145]

R³

reactions

'n,



ő

Bn

• Jayanth [121] • Cu-catalyzed coupling of imines, vinylstannanes, or alkynes and o-bromoaroyl chlorides followed by Pd(0)-catalyzed annulations



Brummond [29]
 tandem cyclopropanation/
Cope rearrangement followed
by a Diels-Alder sequence



Dou [60]
cyclization of thiourea intermediates mediated by low-valent titanium

Dzhavakhishvili [63]
reduction of N2-substituted Gewald's amides with aromatic aldehydes

x

Ghahremanzadeh [77]
4-CR of indolin-2-one, 3-

Bn

ö

 Smith [271]
 Lewis acid catalyzed 3-CR
hetero-Diels-Alder of *N*-arylimines with strained norbornene-derived dienophiles

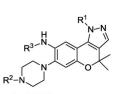


Liu [172]
Ugi and Pictet-Spengler reactions

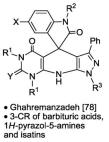


Sotoca [272]
cyclodehydrative 3-CR with 1,3-dicarbonyls





Park [220]regioselective condensation of β -keto aldehydes with hydrazines

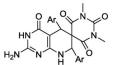




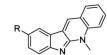
Medimagh [188]
 3-CR cascade process via formation of a transient imine followed by Diels-Alder cycloaddition, oxazolidine ring closure, and lactamization



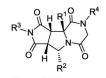
Han [95]
 MW-assisted condensation of 3,5-dibenzylidenepiperidin-4-ones with 2,6-diaminopyridin-4(3*H*)-one and other enamine-like substrates



Jiang [124]
MW-assisted domino synthesis from ArCHO and barbituric acids



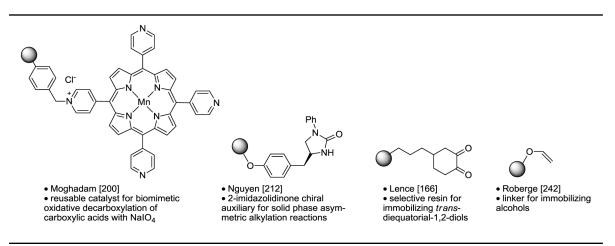
Miller [330]
 reaction of fluorous thiols
 with glyoxamides via
 Pummerer-type cyclization



Werner [328]
 [3+2] cycloaddition of fluorous amino esters, aldehydes, and maleimides

^aAsterisk is the point of attachment to resin.

Polymer-Supported Reagents, Scavengers, and Linkers.



Fluorous Catalysts, Reagents, Scavengers, Linkers and Library Synthesis.

