

Laurea Magistrale in Chimica

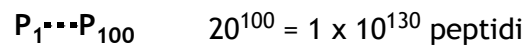
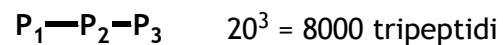
Laboratorio di Chimica Bioorganica

anno accademico 2015 – 2016

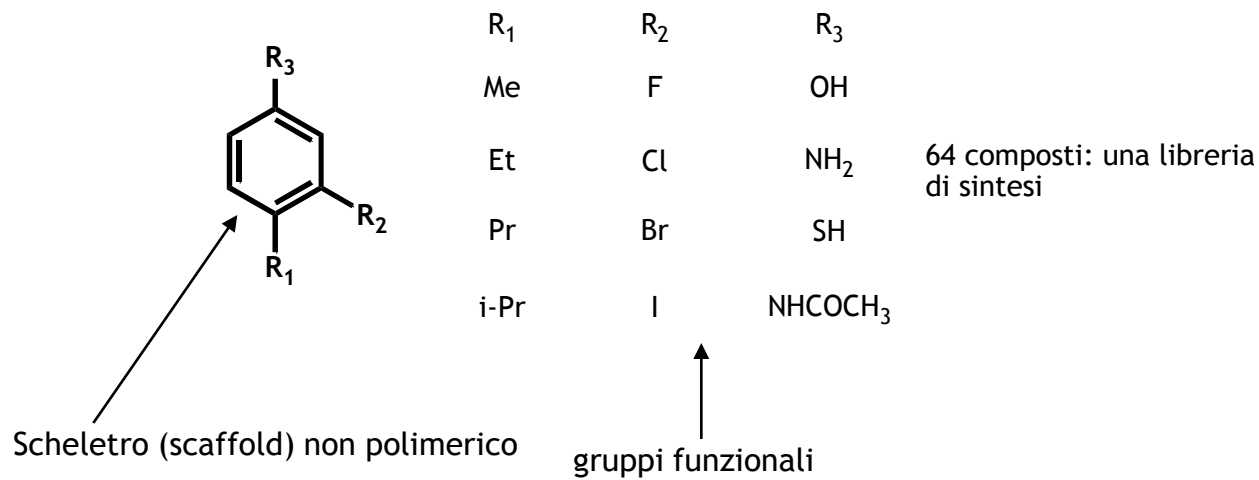
Prima esperienza

Due librerie ortogonali di inibitori della β -secretasi a scaffold diidropirimidinico: sintesi per reazione di Biginelli supportata e valutazione dell'attività biologica.

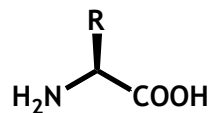
Introduzione alla chimica combinatoriale



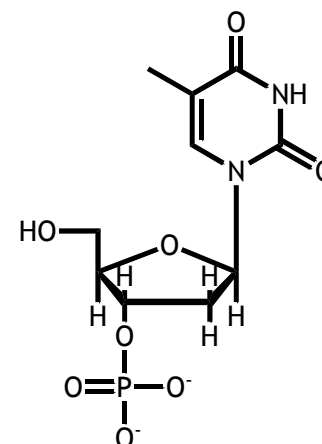
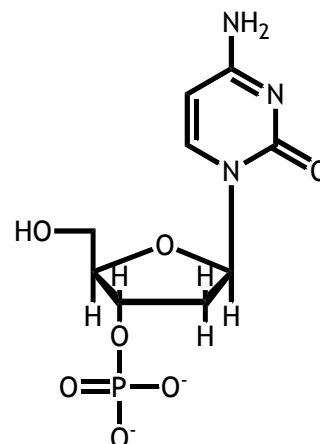
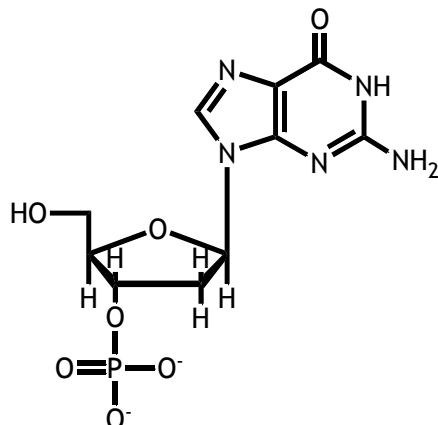
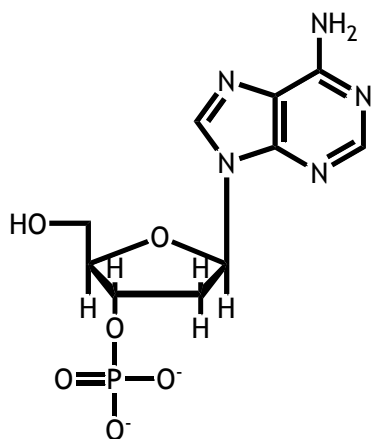
Una libreria di origine biologica, ad es. fagica: $10^6 - 10^8$ membri



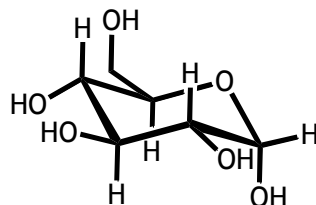
Scheletro polimerico: il gruppo funzionale fa parte del monomero che partecipa alla eteropolimerizzazione.



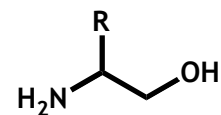
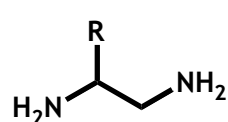
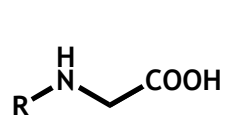
Peptidi: variabilità in R



Acidi nucleici: variabilità in base

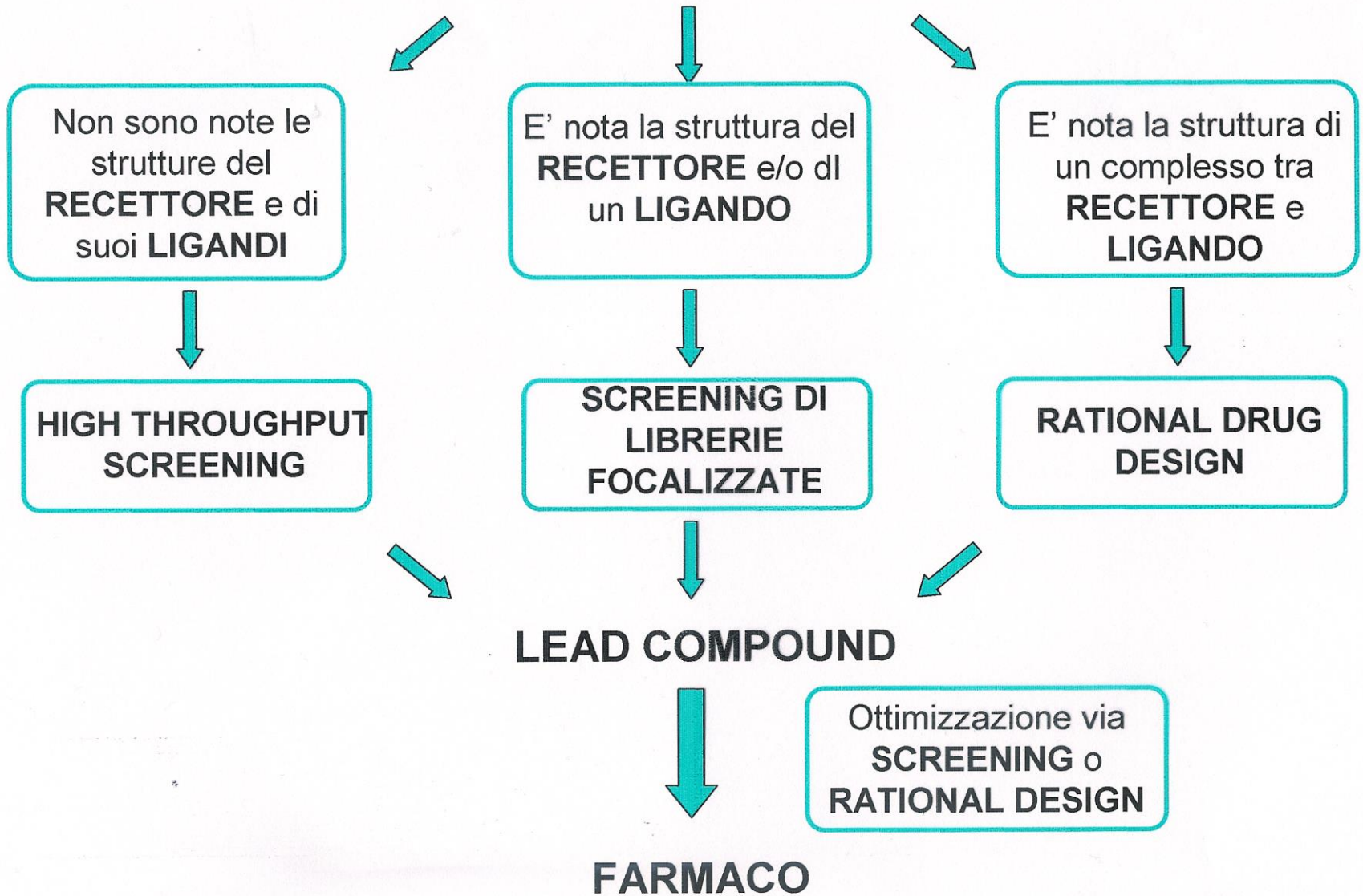


Polisaccaridi: variabilità in struttura, stereochimica, legame glicosidico

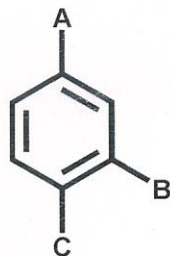


Scheletri polimerici artificiali: peptoidi, poliuree, poliuretani

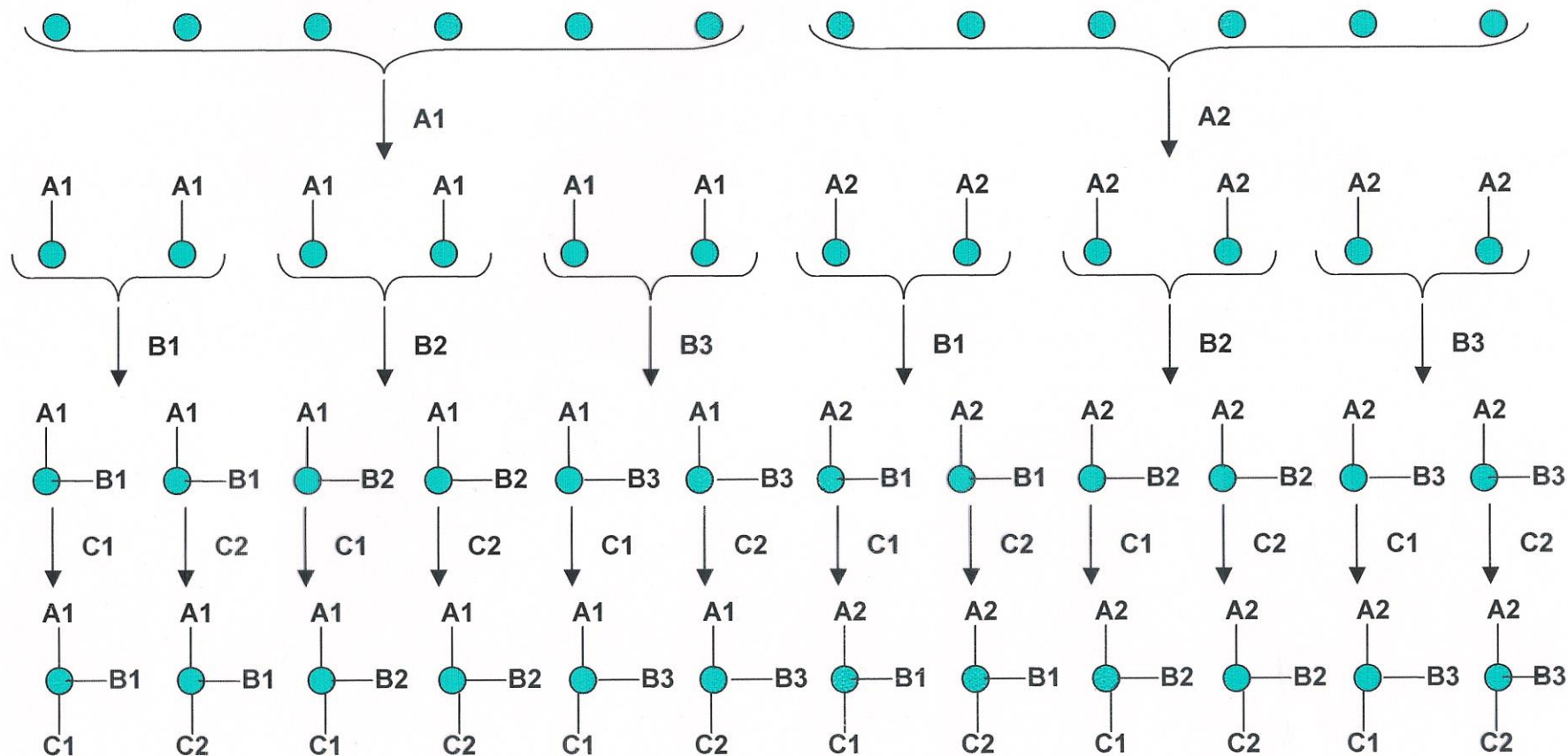
OBIETTIVO



SINTESI PARALLELA DI:

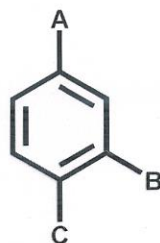


A1, A2 - B1, B2, B3 - C1, C2

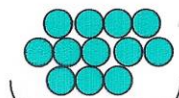


2+6+12=20 reazioni per 12 composti

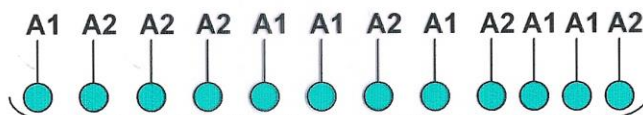
SINTESI IN MISCELA DI:



A1, A2 - B1, B2, B3 - C1, C2



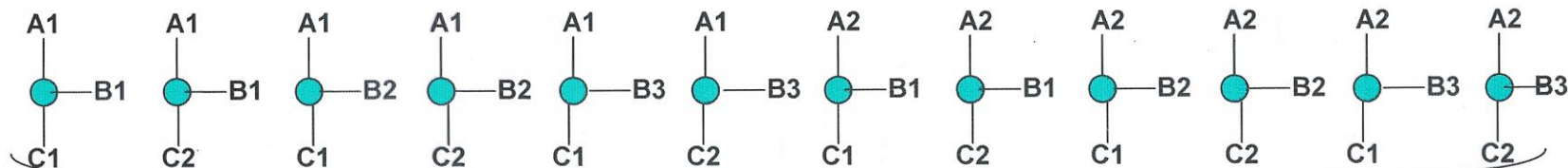
50% A1 + 50% A2



33% B1 + 33% B2 + 33% B3

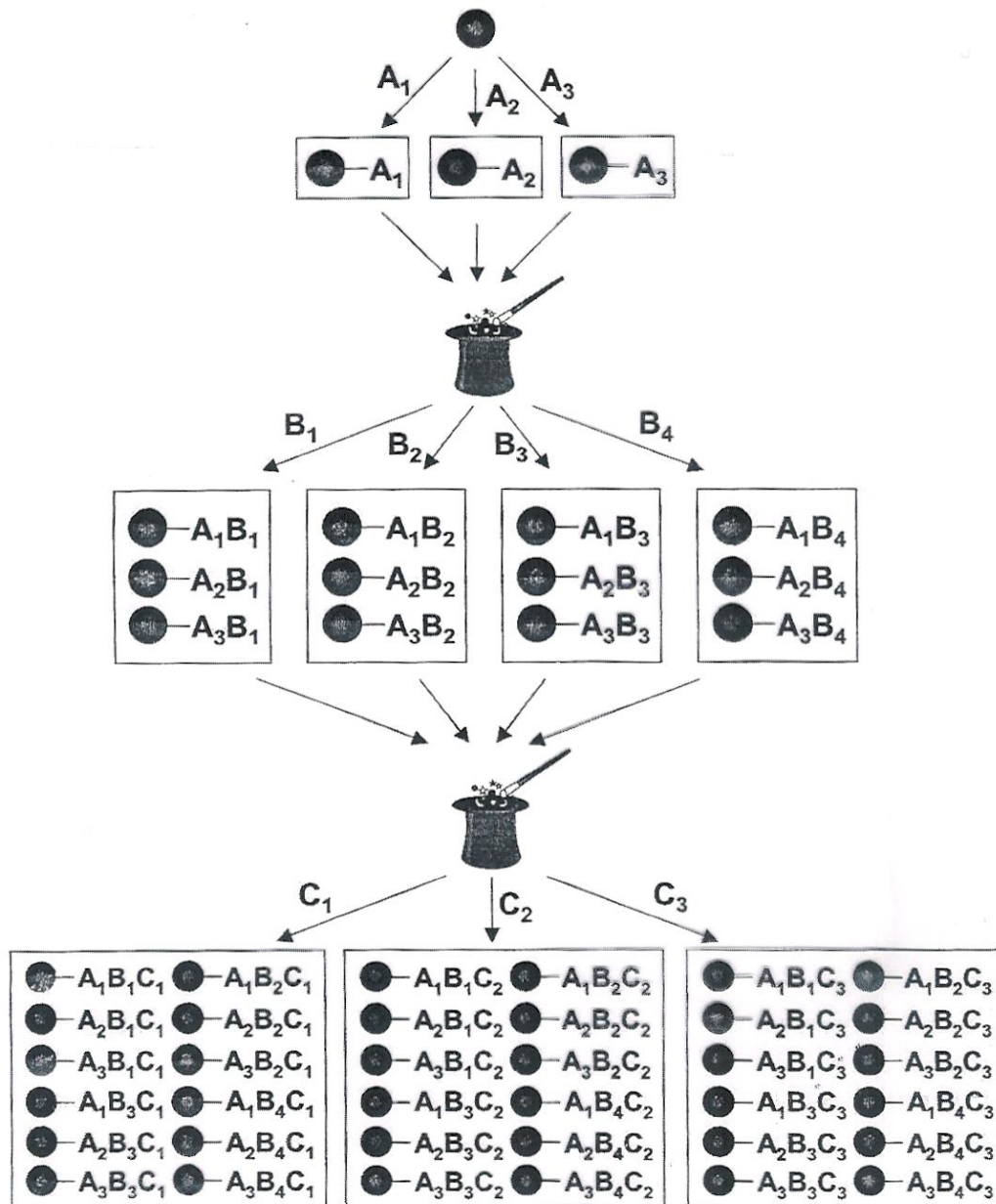


50% C1 + 50% C2



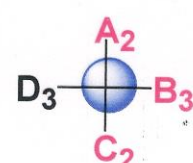
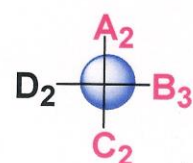
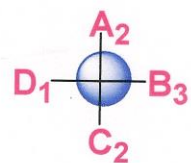
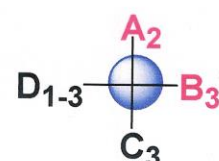
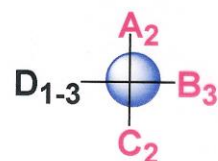
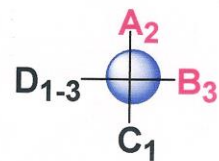
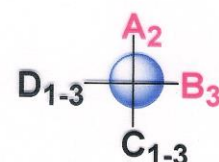
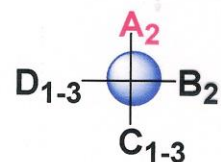
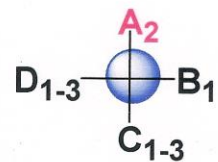
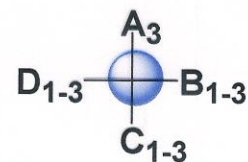
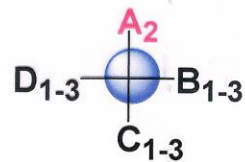
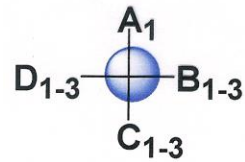
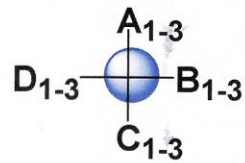
Tre reazioni per una libreria di 12 composti

SPLIT AND COMBINE LIBRARY SYNTHESIS

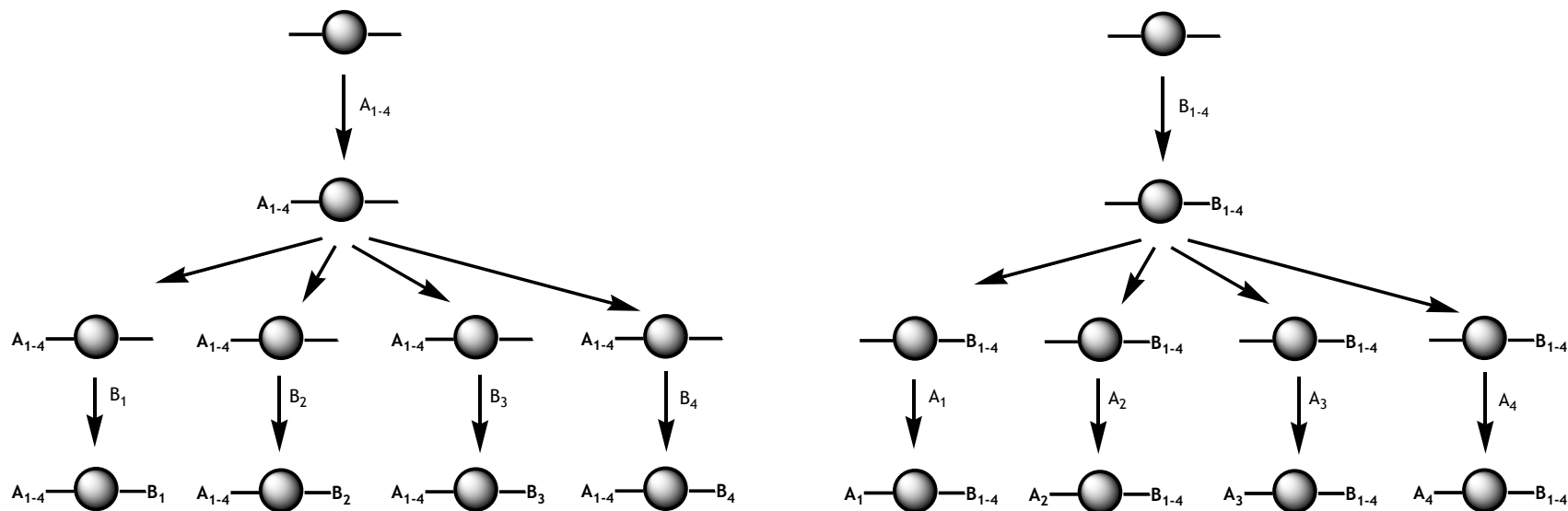


Sintesi in miscela di una libreria composta di una gerarchia di sottolibrerie

DECONVOLUZIONE ITERATIVA

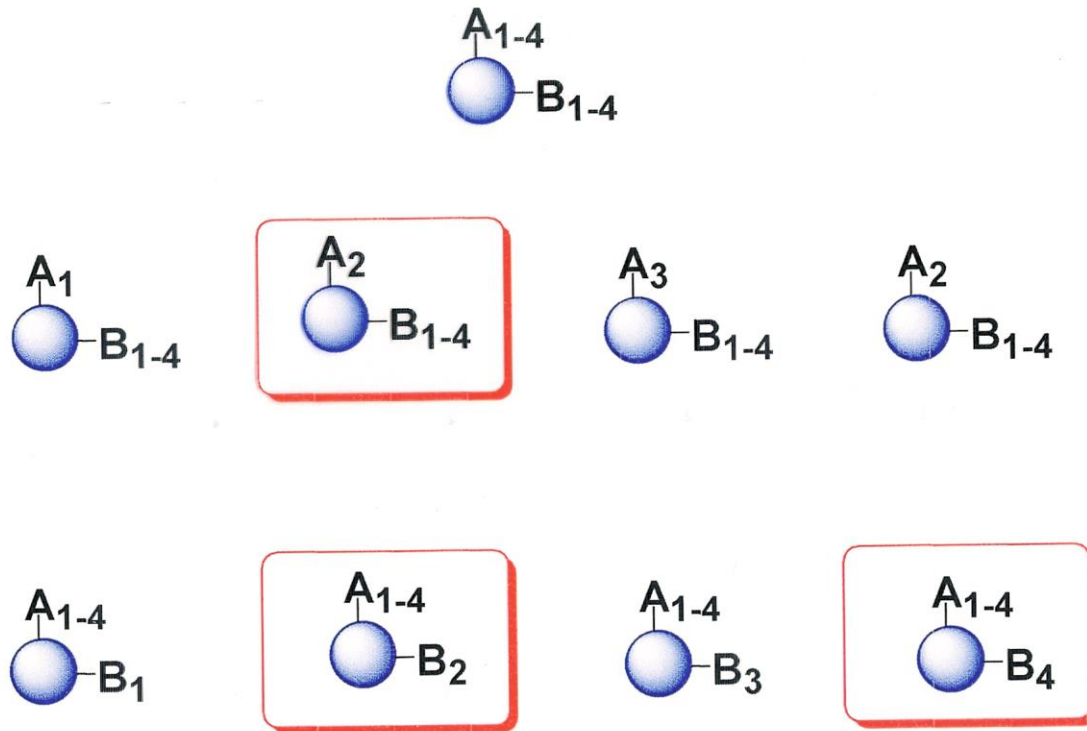


Sintesi in miscela di librerie ortogonali



100 A, 100 B: 10000 composti, 10000 reazioni per la sintesi parallela pura. 202 reazioni in quattro passaggi per la sintesi di librerie ortogonali.

Deconvoluzione di librerie ortogonali



Attivi: A_2B_2 e A_2B_4



La β -secretasi

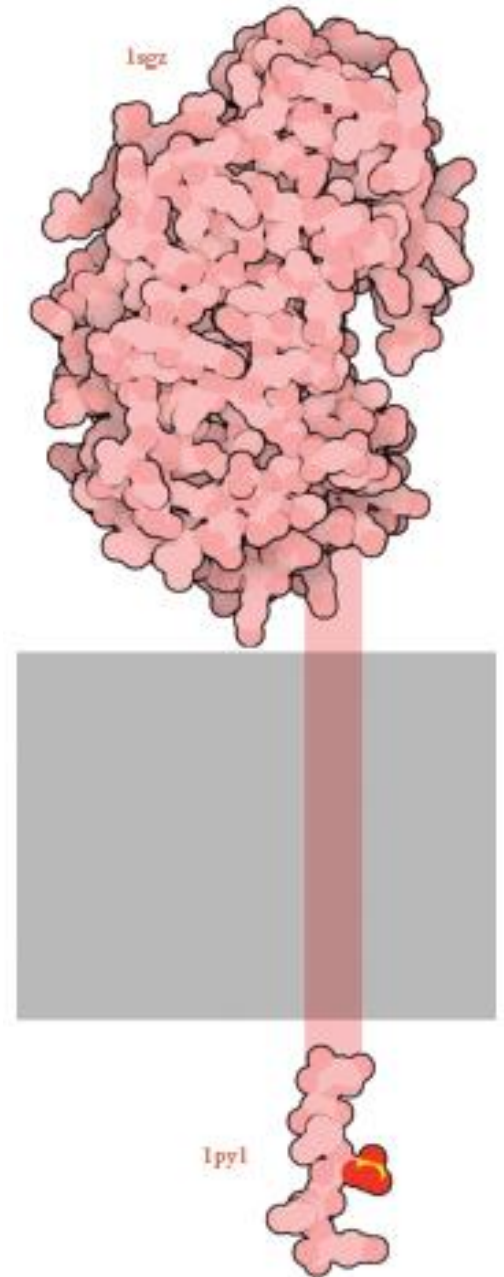
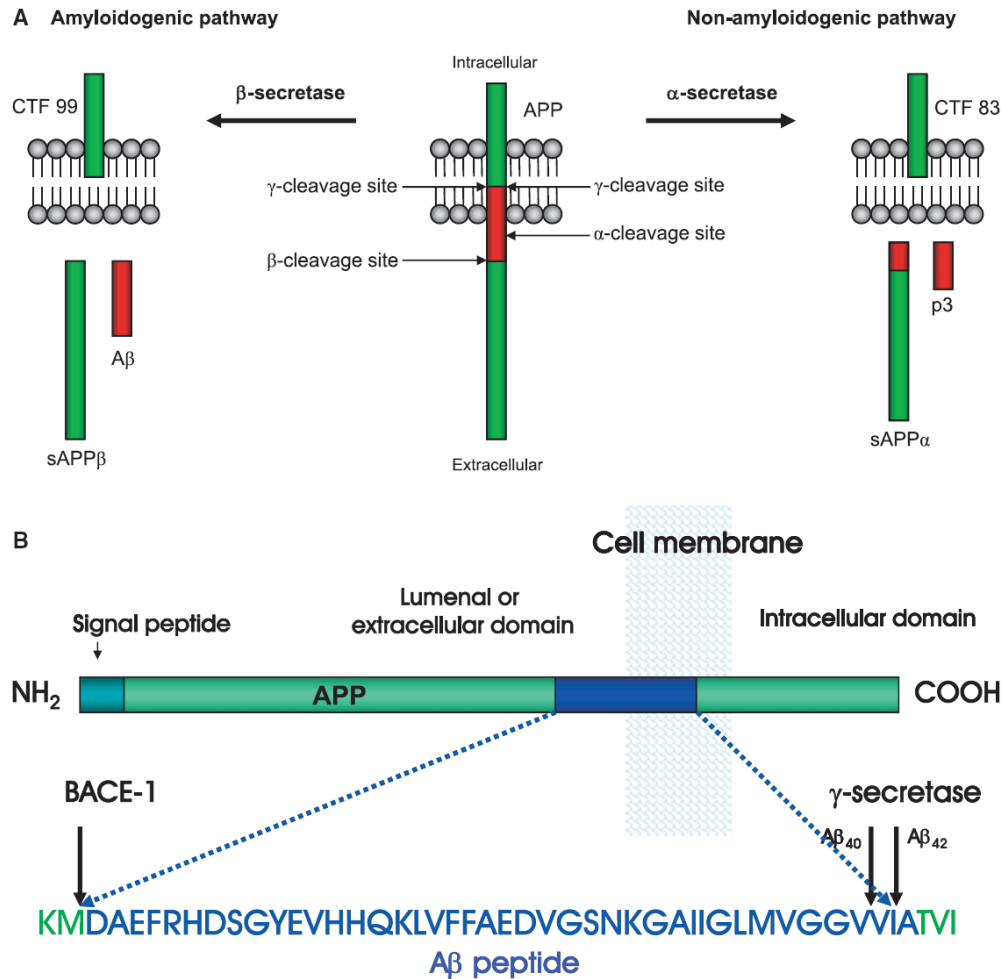
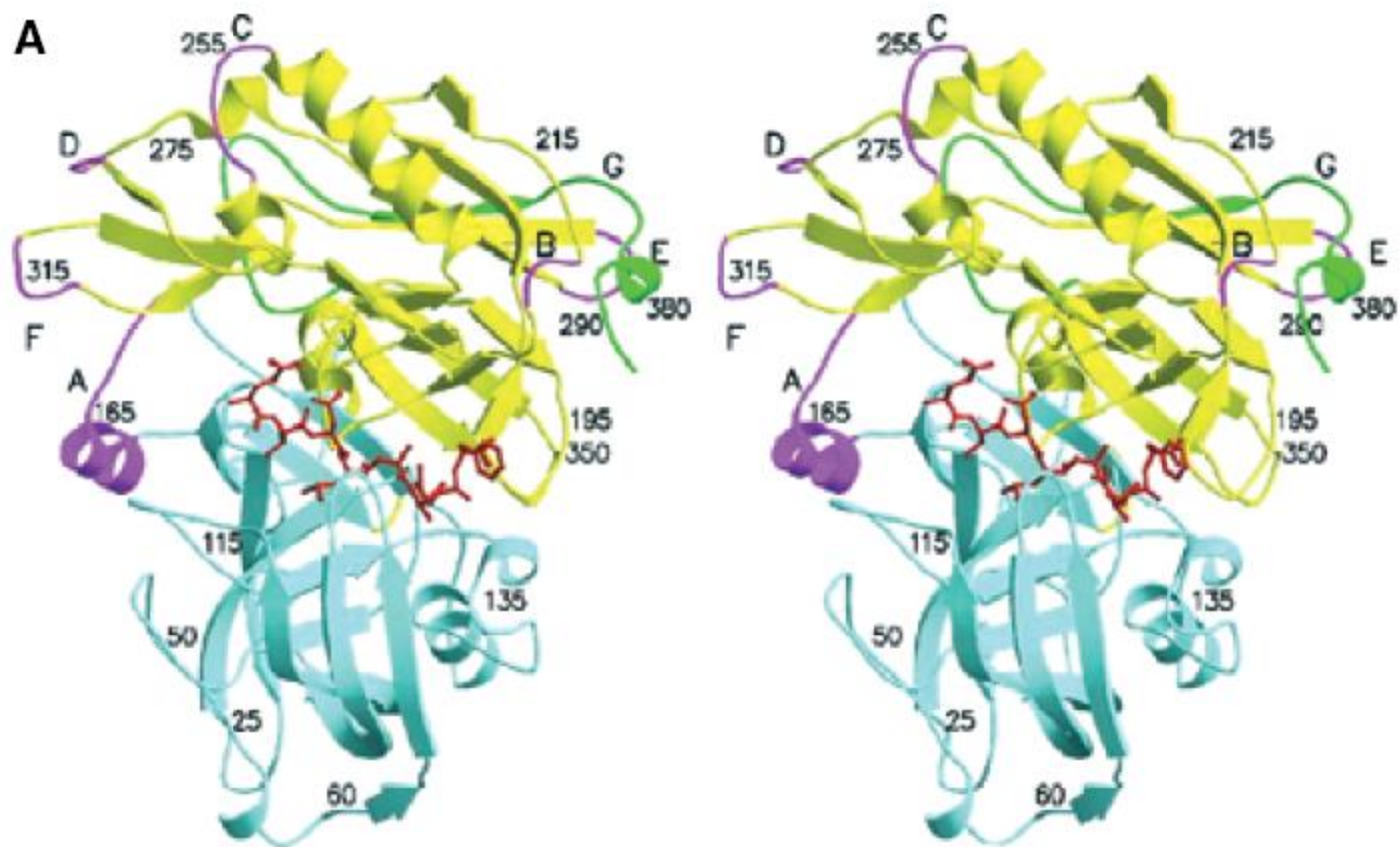
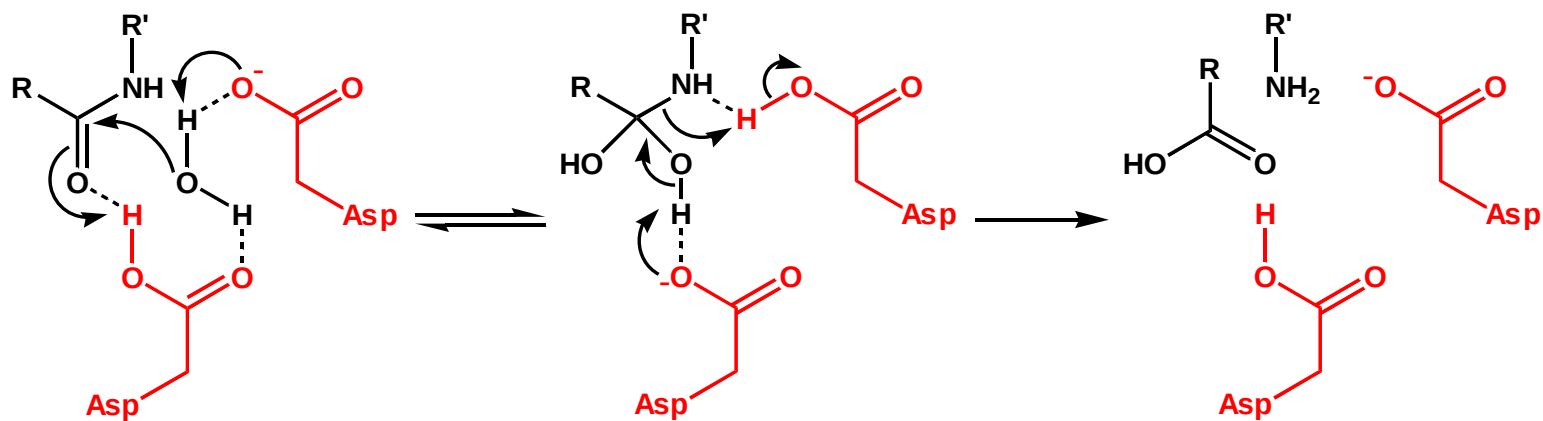
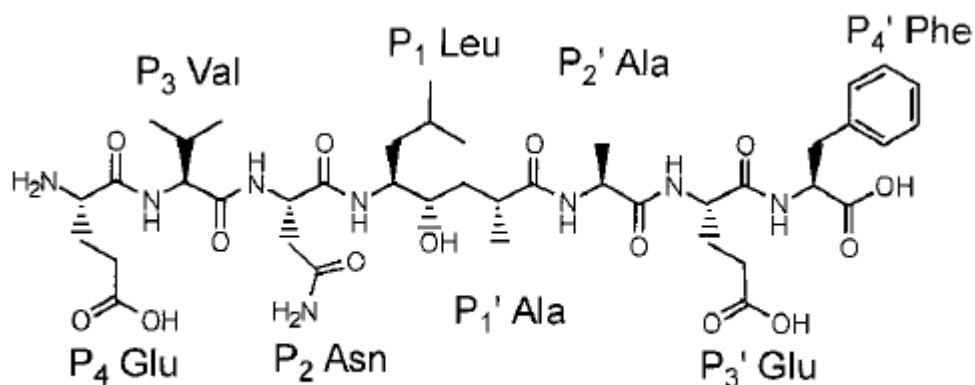


Fig. 1. Processing of APP to form A β peptides. (A) Schematic diagram of the alternative processing pathways of APP. The transmembrane APP undergoes two alternative and competing pathways of metabolism. The major and non-amyloidogenic, or α -secretase, pathway precludes the formation of Alzheimer's A β peptide. The amyloidogenic, or β -secretase, pathway initiates the formation of A β , which is completed by the action of the γ -secretase. α -Secretase has been identified as a zinc metalloproteinase of the ADAMs family, whereas both β - and γ -secretases are membrane-bound aspartic proteinases (see text for full details). (B) Sites of cleavage of APP by β - and γ -secretases to form A β peptides. The sites of the juxtamembrane and intramembrane cleavages of transmembrane APP by β - and γ -secretases, respectively, are indicated by arrows. The γ -secretase cleavages are heterogeneous, mainly producing A β peptides of 40 and 42 amino acids. The amino acid sequences of A β and around the scissile bonds are indicated by the one letter code for amino acids. The sequence shown is the wild-type sequence. The 'Swedish mutant' APP sequence around the β -secretase cleavage site is ...NL/DAEF... rather than ...KM/DAEF... The development of many BACE-1 inhibitors has used the sequence around the scissile bond in the Swedish mutant as the lead for synthetic chemistry to produce potent and selective compounds.

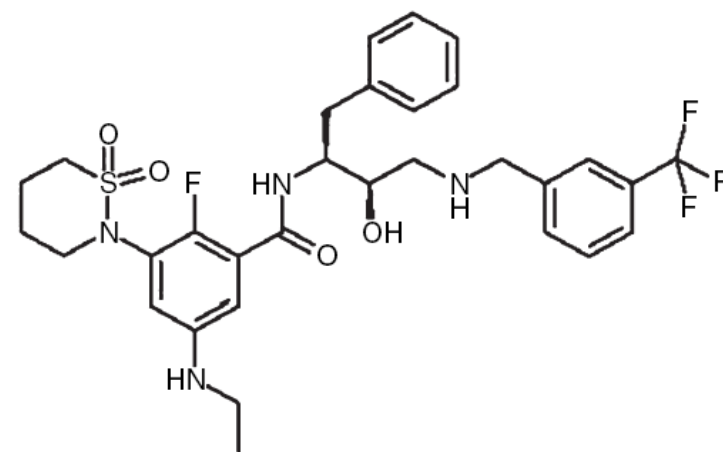


Le proteasi aspartiche

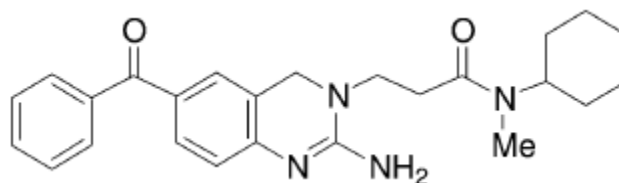




Design of potent inhibitors for human brain memapsin 2 (β -secretase). A. Ghosh et al., J. Am. Chem. Soc. 2000, 122, 3522.



Oral administration of a potent and selective non peptidic BACE-1 inhibitor decreases β -cleavage of amyloid precursor protein and amyloid- β production in vivo. I. Hussain et al. J Neurochem 2007, 100, 802.



2-Amino-3,4-dihydroquinazolines as Inhibitors of BACE-1 (β -Site APP Cleaving Enzyme): Use of Structure Based Design to Convert a Micromolar Hit into a Nanomolar Lead. E. W. Baxter et al., J. Med. Chem. 2006, 50, 4262.

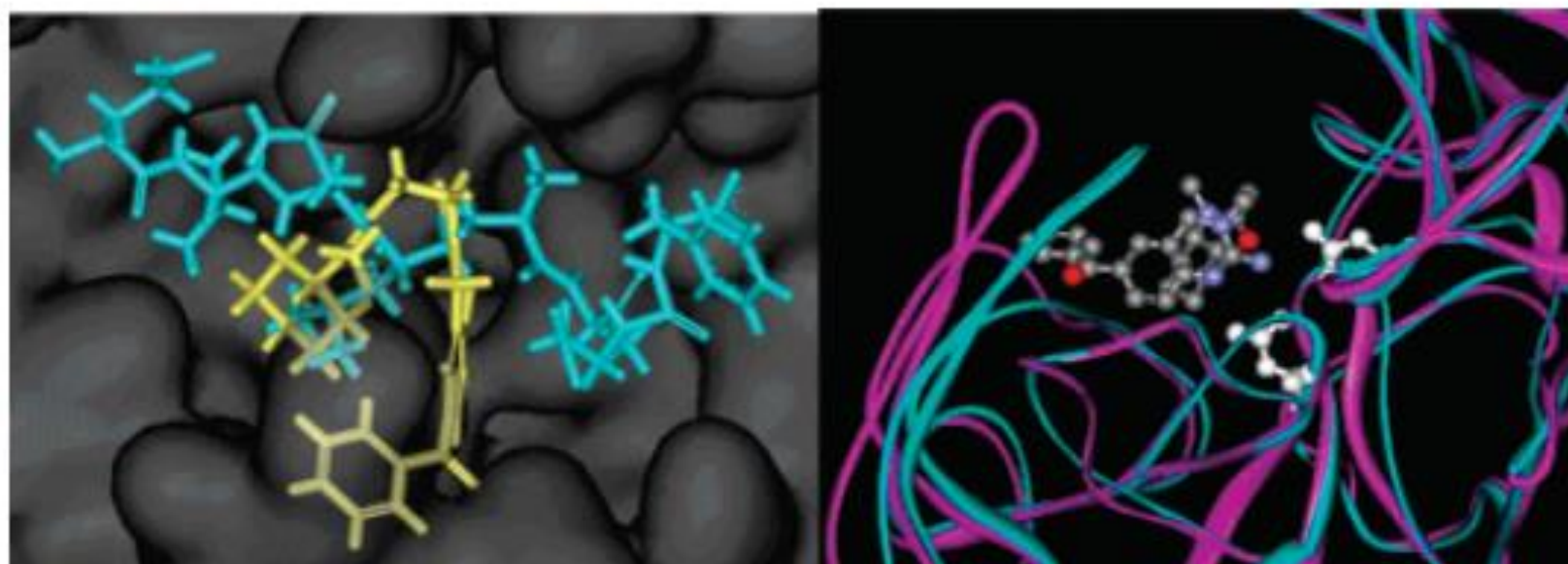


Figure 1. Left: superposition of OM99-2 (peptidic inhibitor) and **1** in the active site of BACE-1. Right: ribbon diagram representation of **1** with the two catalytic aspartic acids highlighted bound into BACE-1 (maroon), showing the flap over the active site moved when compared with BACE-1 when OM99-2 is bound in (turquoise).

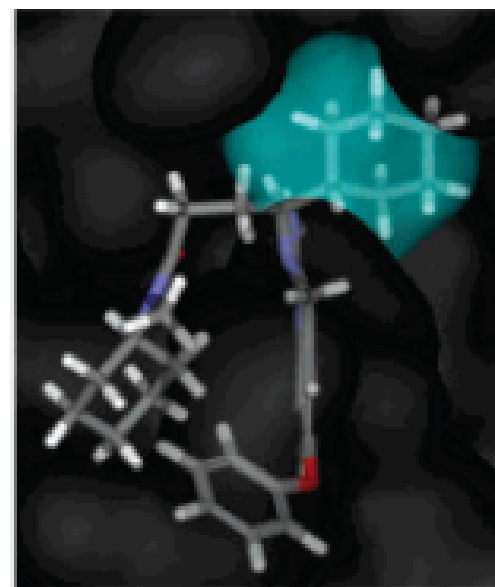
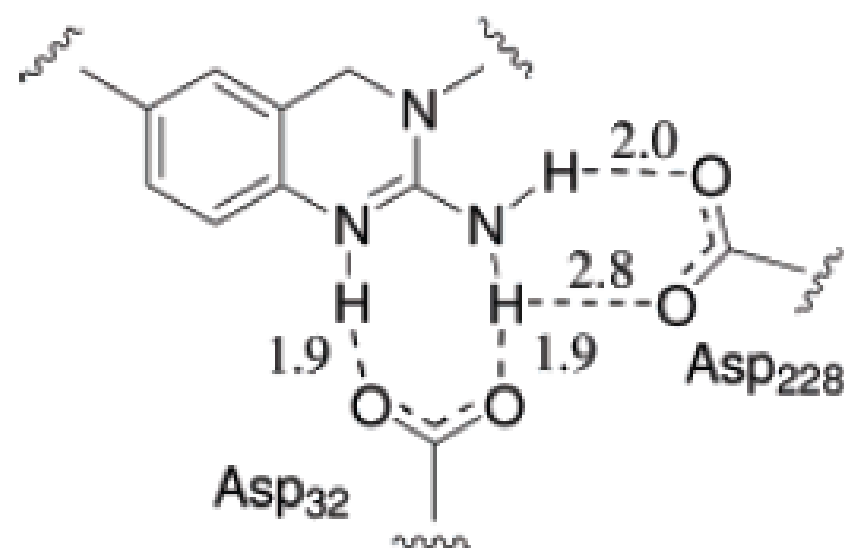


Figure 2. Left: the 2-amino-3,4-dihydroquinazoline fragment of **1** and the Asp₃₂ and Asp₂₂₈ sidechains of BACE-1, with H-bond distances in Å between the Asp carboxylate oxygens and the hydrogens. Right: the X-ray structure of **3a** bound in to BACE-1, showing the α -cyclohexyl substituent filling the S₁' pocket.

La reazione multicomponente di Biginelli

Biginelli P. Derivati aldeiduredici degli eteri acetil- e dossal-acetico,
Gazzetta Chimica Italiana, **1893**, 23, 360 – 416

QSAR
& **Combinatorial Science**

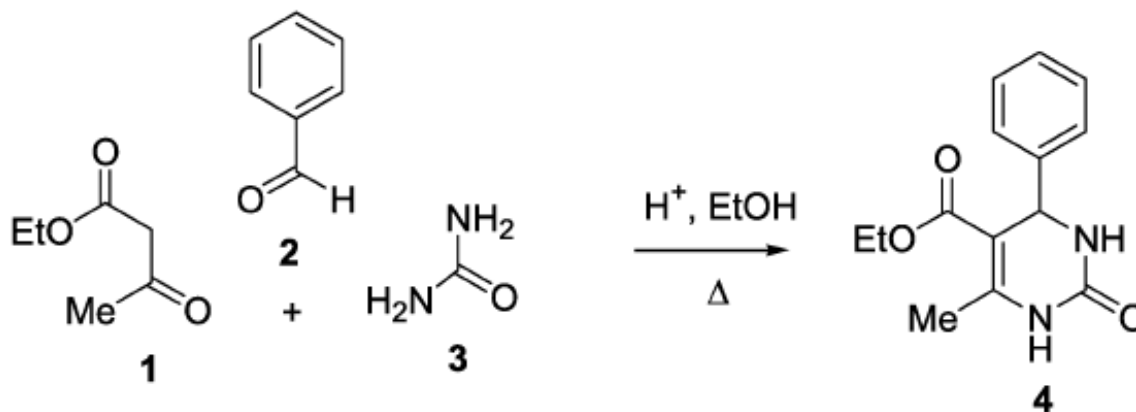
C. Oliver Kappe

The Generation of Dihydropyrimidine Libraries Utilizing Biginelli Multicomponent Chemistry

C. Oliver Kappe

Institute of Chemistry, Karl-Franzens-University Graz, Heinrichstrasse 28, A-8010 Graz (Austria), Fax: (+43)316-380-9840,
E-mail: oliver.kappe@uni-graz.at

QSAR Comb. Sci. 22 (2003) DOI: 10.1002/qsar.200320001



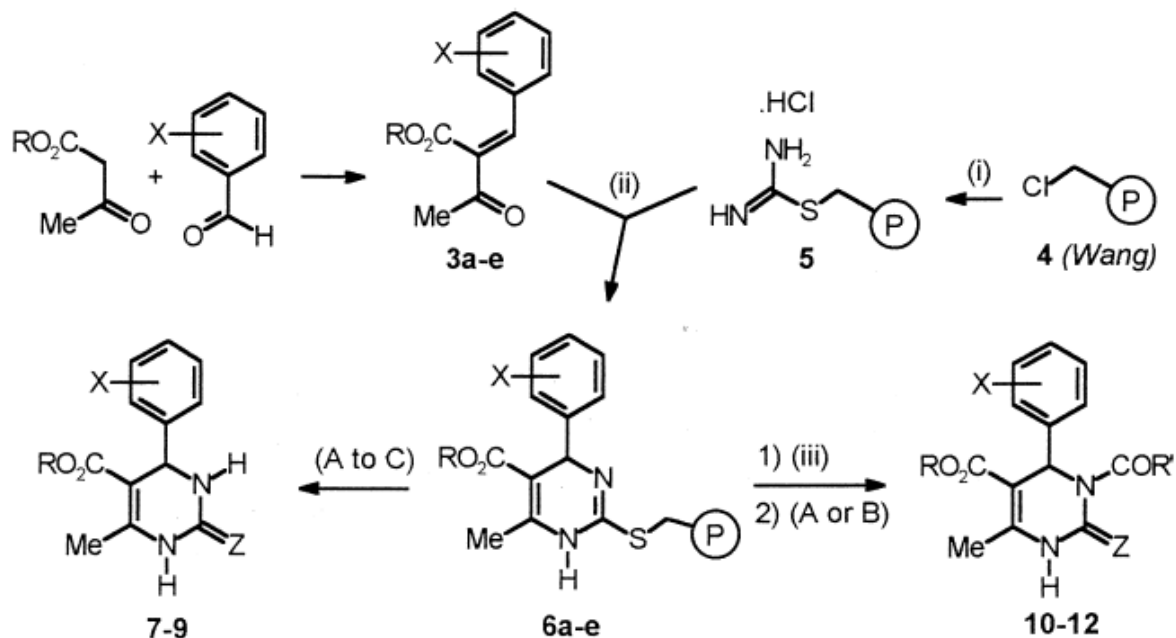
Scheme 1. The original Biginelli dihydropyrimidine synthesis (1893).



Highly Versatile Solid Phase Synthesis of Biofunctional 4-Aryl-3,4-dihydropyrimidines Using Resin-Bound Isothiourea Building Blocks and Multidirectional Resin Cleavage[†]

C. Oliver Kappe*

Institute of Organic Chemistry, Karl-Franzens-University Graz, Heinrichstrasse 28, A-8010 Graz, Austria



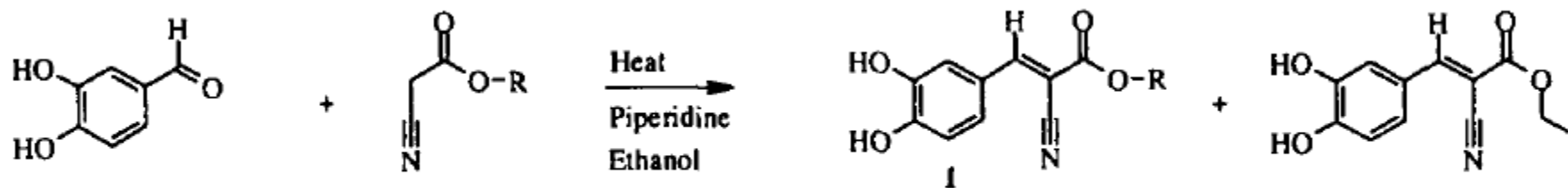
Scheme 2. Reagents and conditions:²⁴ (i) thiourea, NMP, 75 °C, 16 h; (ii) NMP, Cs_2CO_3 , 90 °C, 16 h; (iii) $\text{Cl}-\text{COR}'$, pyridine, CH_2Cl_2 , 0 °C→rt, 6 h; (A) dioxane, ethanol, AcOH, H_2O , reflux, 16 h; (B) TFA, EtSH, CH_2Cl_2 , rt, 16 h; (C) dioxane, MeCN, NH_4OAc , reflux, 8 h.

New solid phase Knoevenagel catalyst

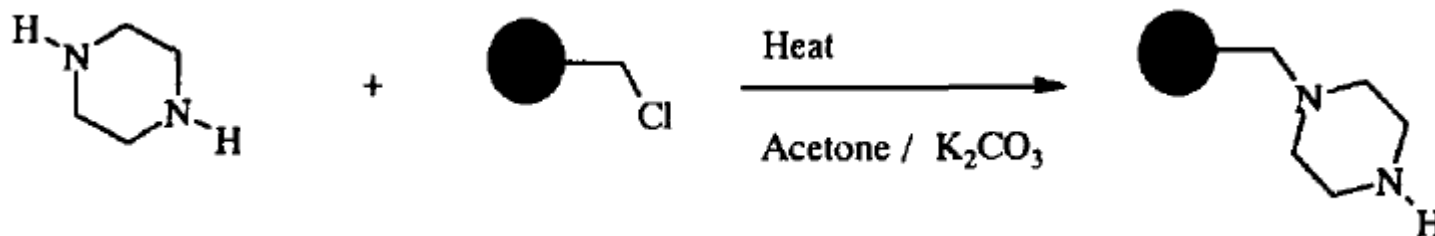
Julie Simpson,* Daniel L. Rathbone and David C. Billington

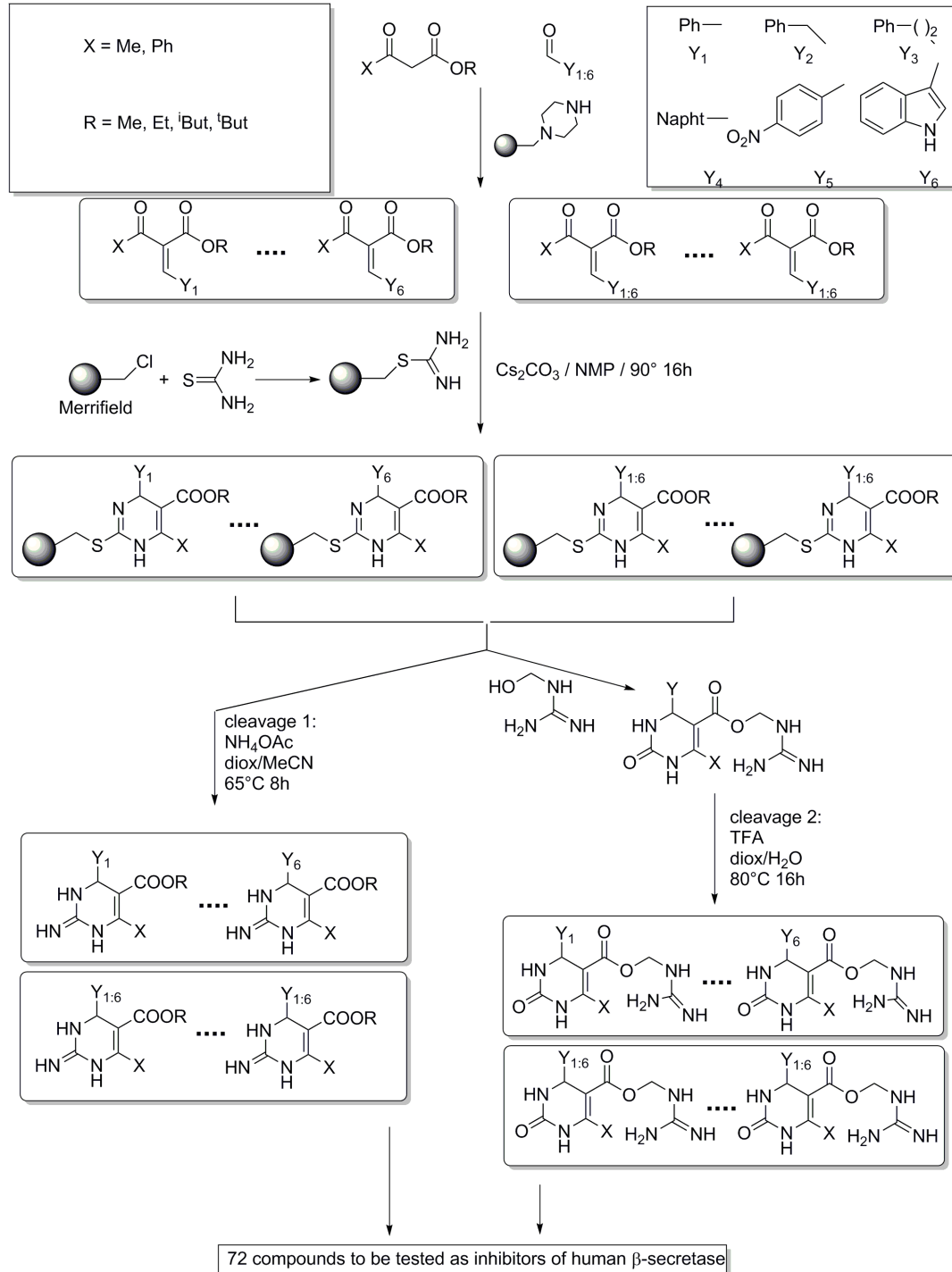
Pharmaceutical Sciences Research Institute, Aston University, Birmingham, B4 7ET, UK

Received 27 May 1999; revised 22 July 1999; accepted 27 July 1999



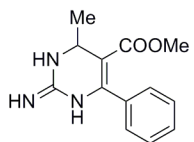
R = Simple and functionalised alkyl groups





X1

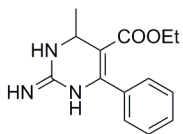
Y1



Mol. Wt.: 245.3

X2

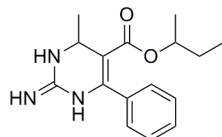
Y2



Mol. Wt.: 259.3

X3

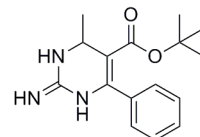
Y3



Mol. Wt.: 287.4

X4

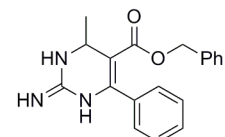
Y4



Mol. Wt.: 287.4

X5

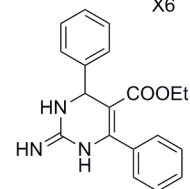
Y5



Mol. Wt.: 321.4

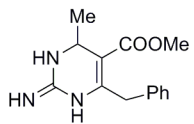
X6

Y6



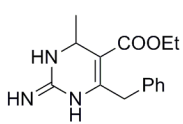
Mol. Wt.: 321.4

Y1



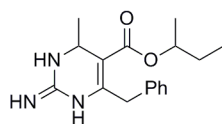
Mol. Wt.: 259.3

Y2



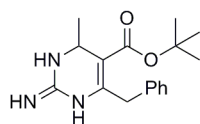
Mol. Wt.: 273.3

Y3



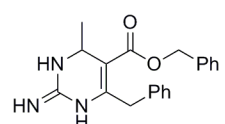
Mol. Wt.: 301.4

Y4



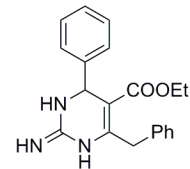
Mol. Wt.: 301.4

Y5



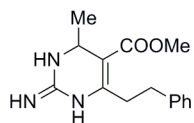
Mol. Wt.: 335.4

Y6



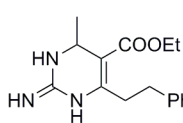
Mol. Wt.: 335.4

Y1



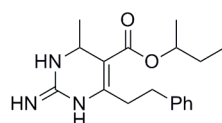
Mol. Wt.: 273.3

Y2



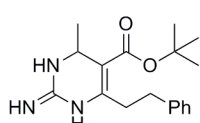
Mol. Wt.: 287.4

Y3



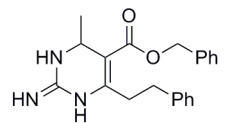
Mol. Wt.: 315.4

Y4



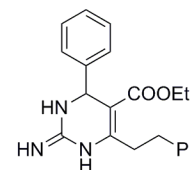
Mol. Wt.: 315.4

Y5



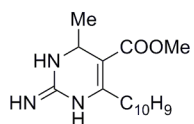
Mol. Wt.: 349.4

Y6



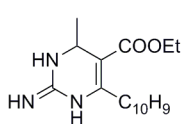
Mol. Wt.: 349.4

Y1



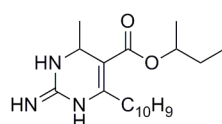
Mol. Wt.: 295.3

Y2



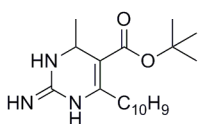
Mol. Wt.: 309.4

Y3



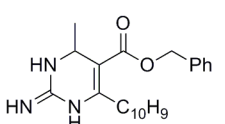
Mol. Wt.: 337.4

Y4



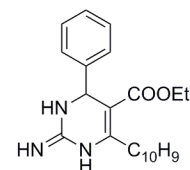
Mol. Wt.: 337.4

Y5



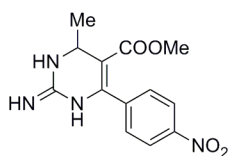
Mol. Wt.: 371.4

Y6



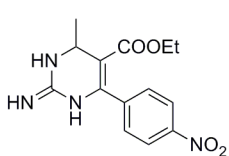
Mol. Wt.: 371.4

Y1



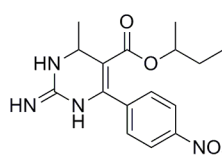
Mol. Wt.: 290.3

Y2



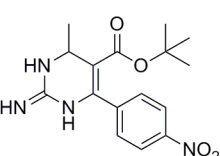
Mol. Wt.: 304.3

Y3



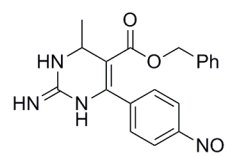
Mol. Wt.: 332.4

Y4



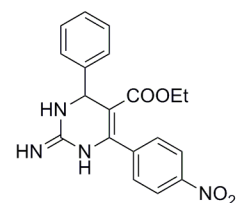
Mol. Wt.: 332.4

Y5



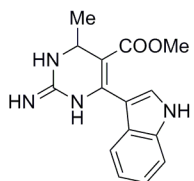
Mol. Wt.: 366.4

Y6



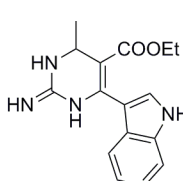
Mol. Wt.: 366.4

Y1



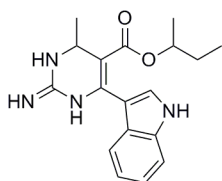
Mol. Wt.: 284.3

Y2



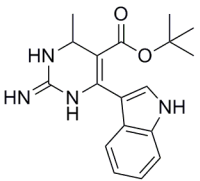
Mol. Wt.: 298.3

Y3



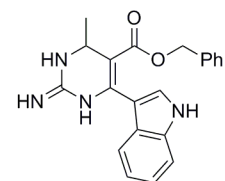
Mol. Wt.: 326.4

Y4



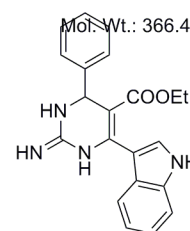
Mol. Wt.: 326.4

Y5



Mol. Wt.: 360.4

Y6



Mol. Wt.: 360.4