

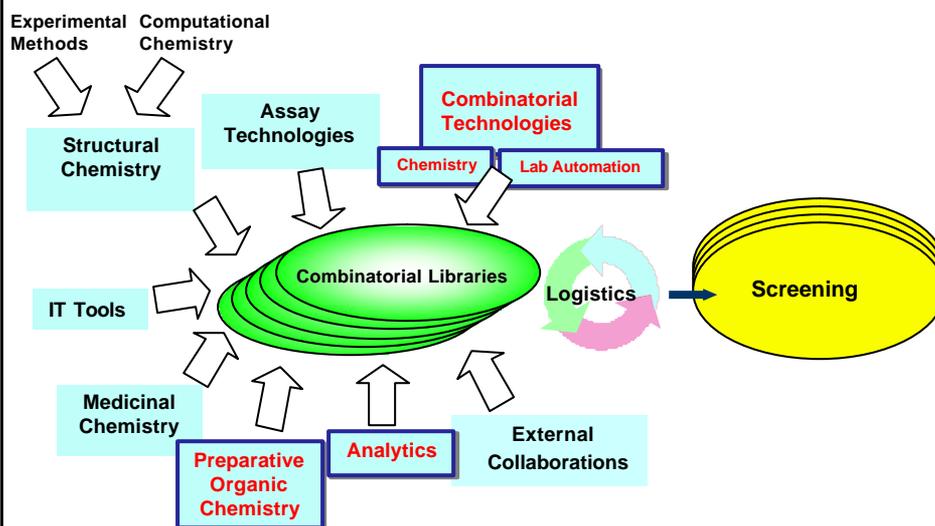
High-throughput synthesis on solid phase

Towards rigorous quality standards and quantitative characterization without compromising efficiency

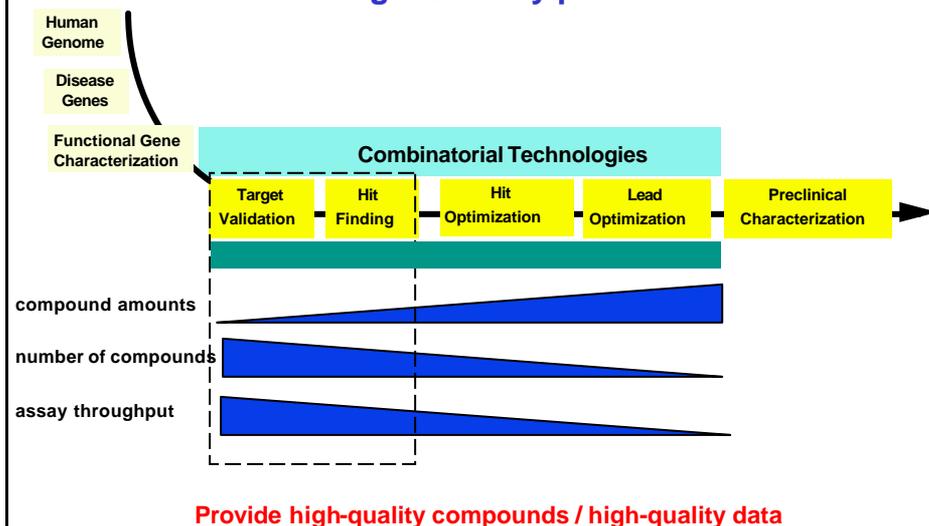
Eduard Felder *et al.*

Dept. of Chemistry - Discovery Research - Pharmacia Italia

Network of Activities



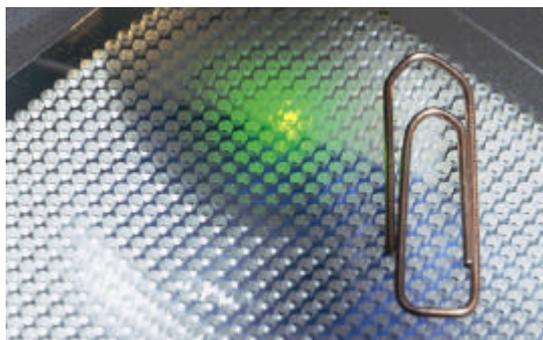
Drug discovery process



Assay Sensitivity and Throughput

UHTS Plate: > 2000 wells / plate
Assay Volume: 1-2 μ l

Microtiter Plate: 96 wells / plate
Assay Volume: 200 μ l



Process Development Areas

LEAD FINDING

LEAD OPTIMIZATION

Hit Generation

Hit Optimization

Parallel Medicinal Chemistry

Avenues:

'Brute Force'

- compd # - - (combinatorial)
- screening data - - (Ultra-HTS)

Screening by NMR
(novel scaffolds)

Combinatorial target-guided
ligand assembly

Technical Development Areas:

- Synthesis ® Automation
 Separations ® High resolution / throughput
 MS-directed fraction collection
 Logistics
 Analytics ® Process control analytics in
 'new formats':
 - on solid phase: IR / NMR
 - in microtiterplates: LC-MS;
 NMR, quant. N-detection

Chemistry Progress: Solid Phase

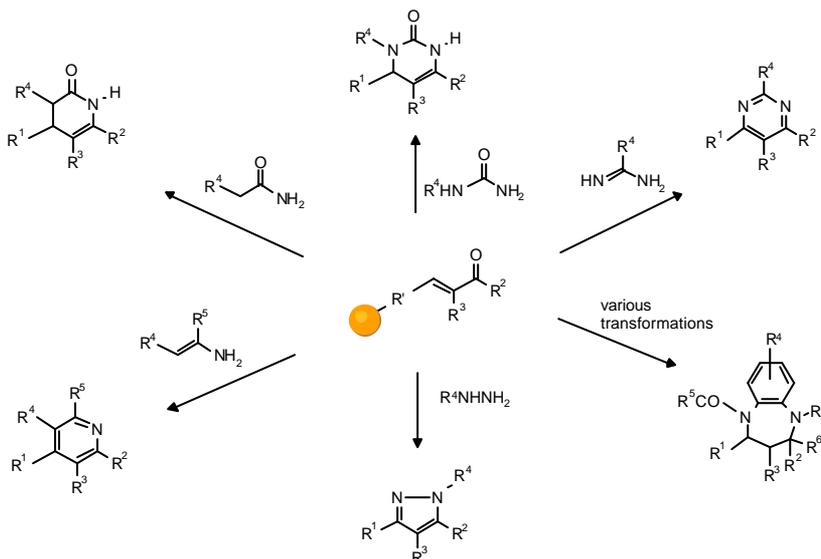
E. Felder - Combinatorial Chemistry

September 2002

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Diversity Platform

NOVARTIS



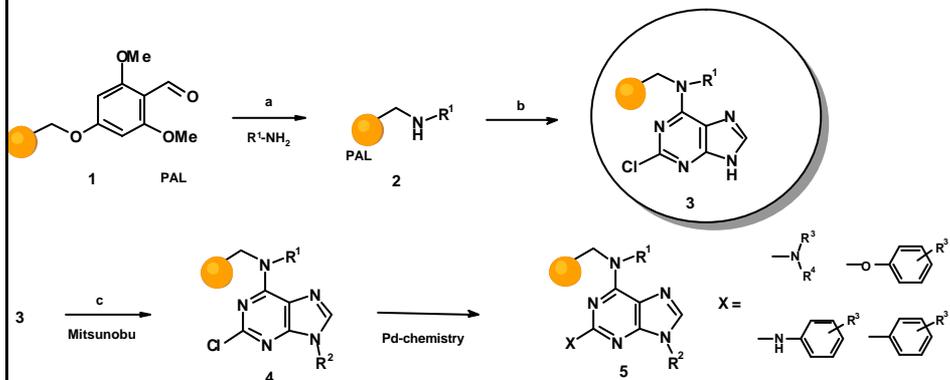
A.L. Marzinzik and E.R. Felder, *J. Org. Chem.* **63**, 723 (1998).

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Derivatizing solid supported core scaffolds with common reactions

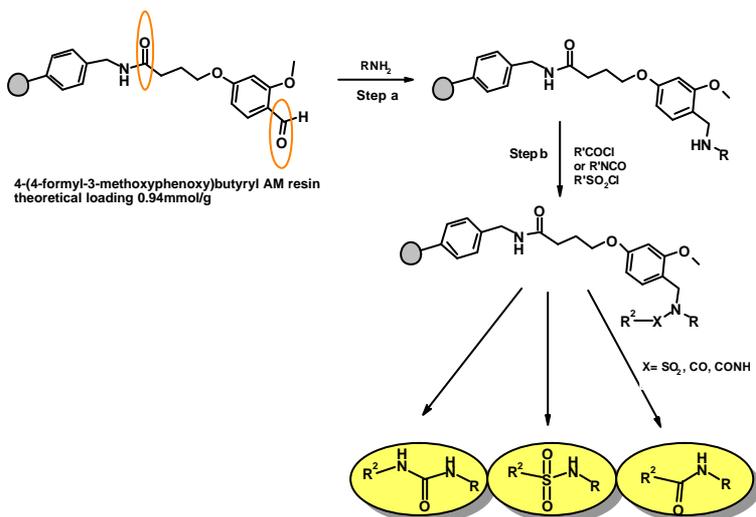


S. Ding et al. *J. Am. Chem. Soc.* **124**, 1594 (2002).

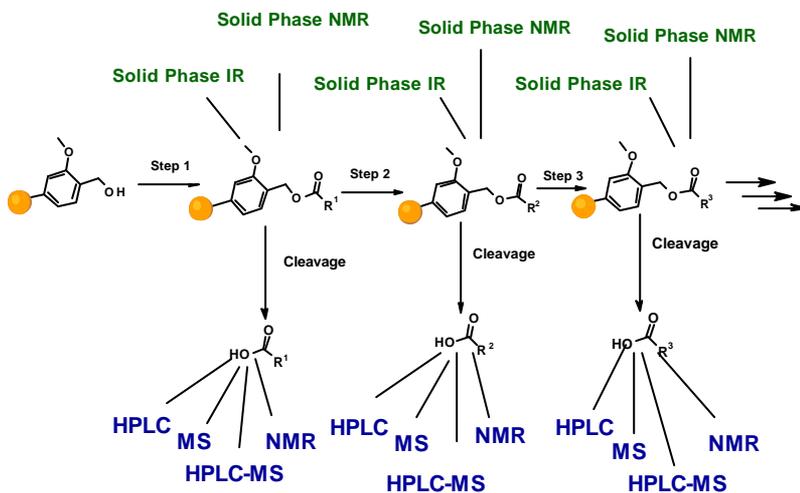
Solid Phase Synthesis - Process Control

- Single Bead IR
- Magic Angle Spinning NMR
- Quantitation by NMR
- Direct-Injection NMR for High-Throughput Spectroscopy
- LC-MS
- HPLC Online Nitrogen Detection

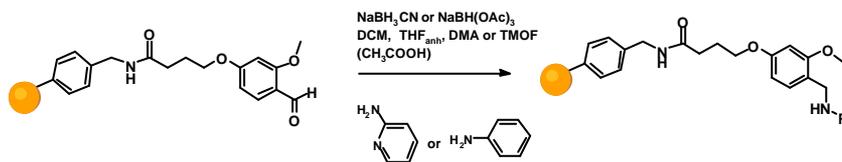
Synthesis Pathway (Example)



Solid Phase Synthesis: Monitoring progress



Reductive Amination - Feasibility Study



STARTING RESIN IR

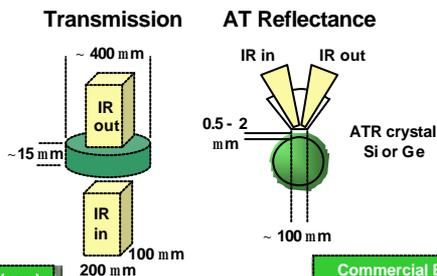
Different analytical techniques were compared

solid phase IR,
MAS NMR,
HPLC

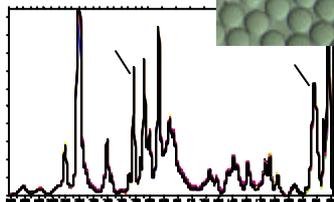
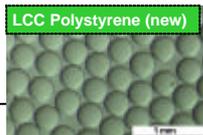
On-bead IR



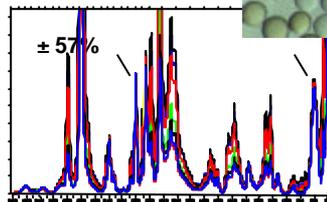
The uniformity of bead loading is measured with microscope IR spectroscopy. The information obtained is unaffected by cleavage efficiency and post-cleavage handling.



Homogeneous bead population

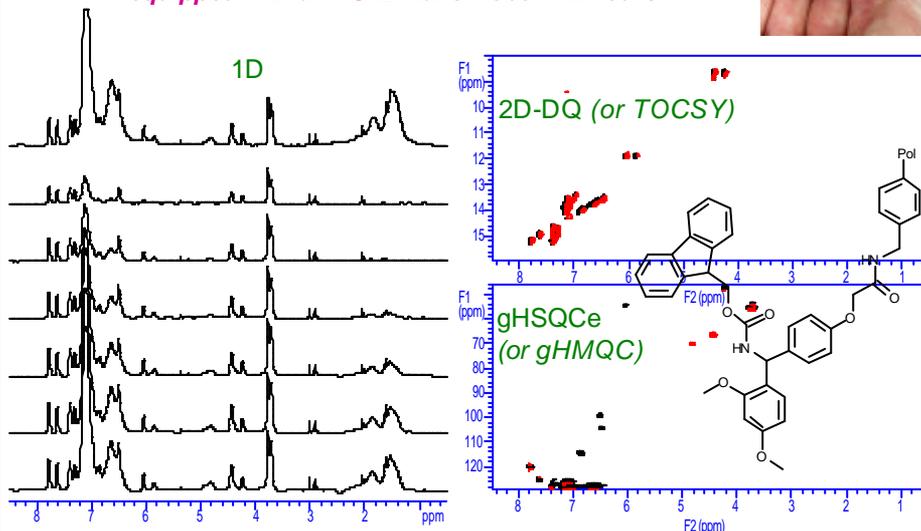


Heterogeneous commercial batch

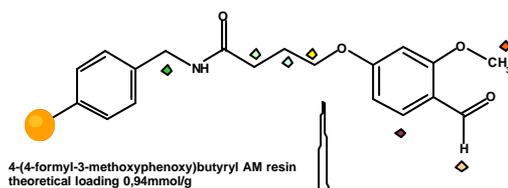


NMR: Our Procedure for SPS-bound cpds

1-3 mg of resin in 40 ml of CD_2Cl_2 , Inova-500 equipped with a PFG-ID NanoProbe: 1-2 hours

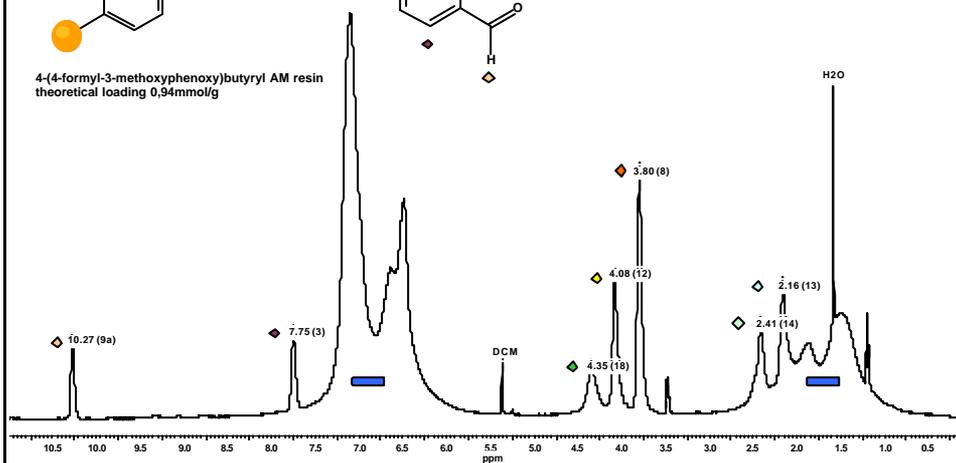


1H MAS NMR Spectra of the formyl resin



Instrument VARIAN INOVA 500
nano probe 40 μ l

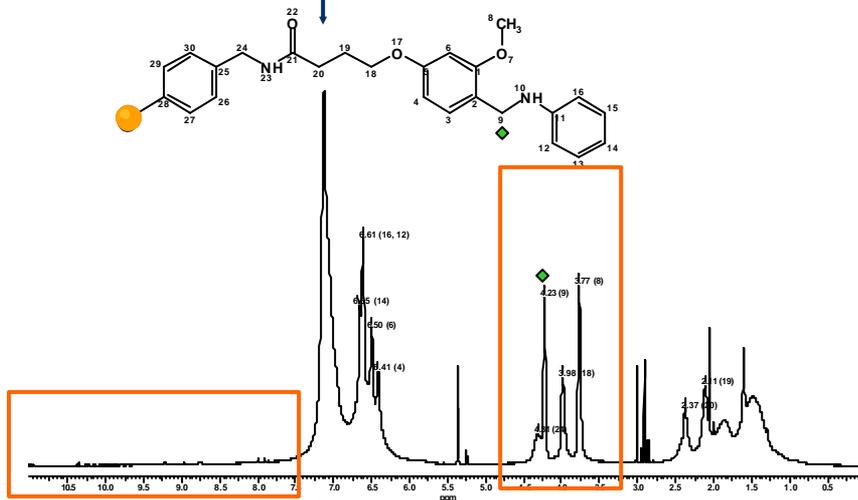
1-3 mg of resin swollen in CD_2Cl_2
scan # 16 t=10ms 3 min exp



Polymeric matrix: Broad signals

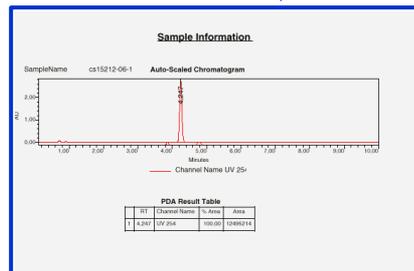
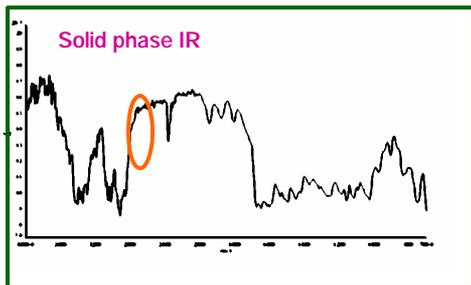
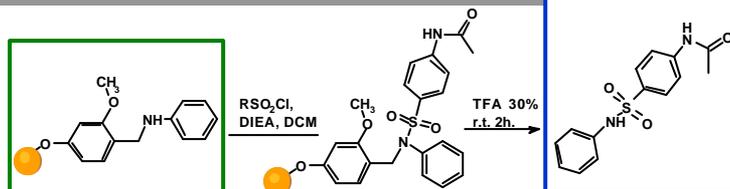
Aniline derivative

Reaction one pot
 aniline 5eq., CH_3COOH 1 eq.,
 $\text{NaBH}(\text{OAc})_3$ 3 eq., in THF o.n.



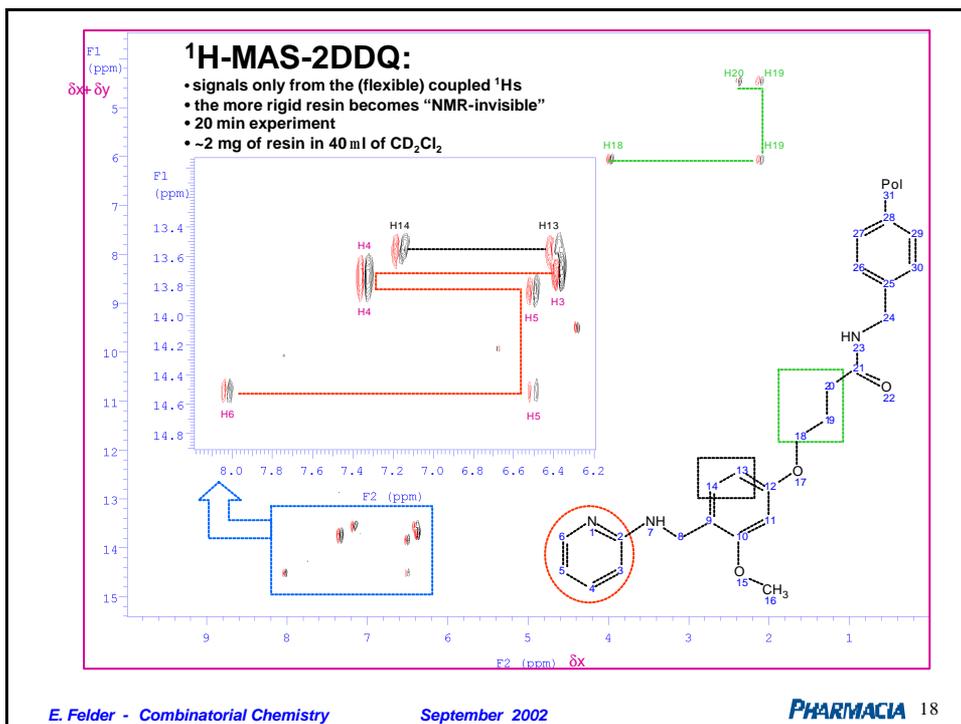
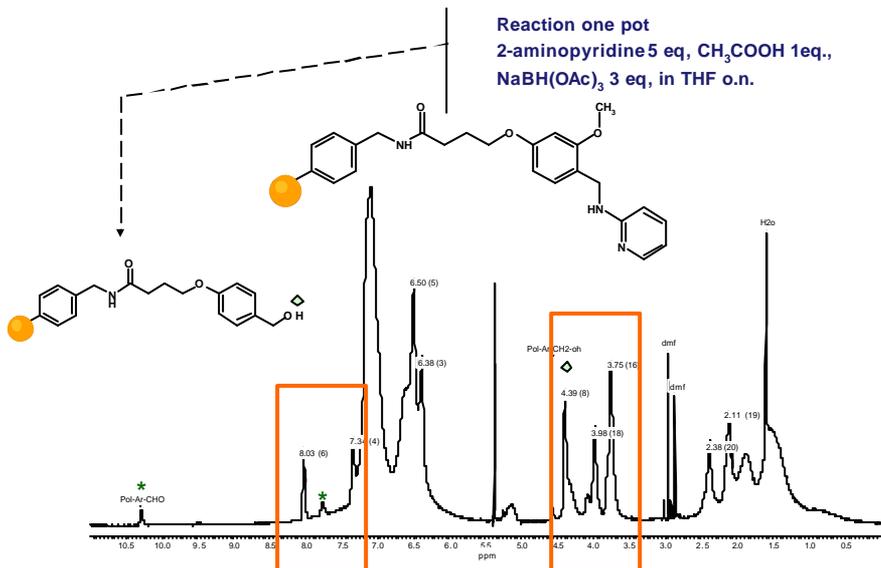
Aniline derivative: Solid phase IR and HPLC after cleavage

aniline,
 CH_3COOH , $\text{NaBH}(\text{OAc})_3$
 THF o.n.

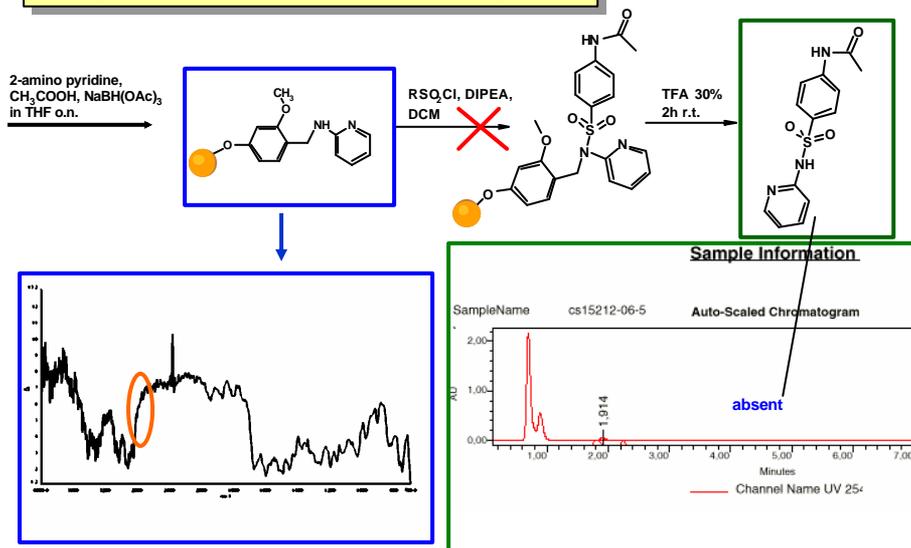


HPLC @254 A= 12495214 after reaction with
 RSO_2Cl and cleavage

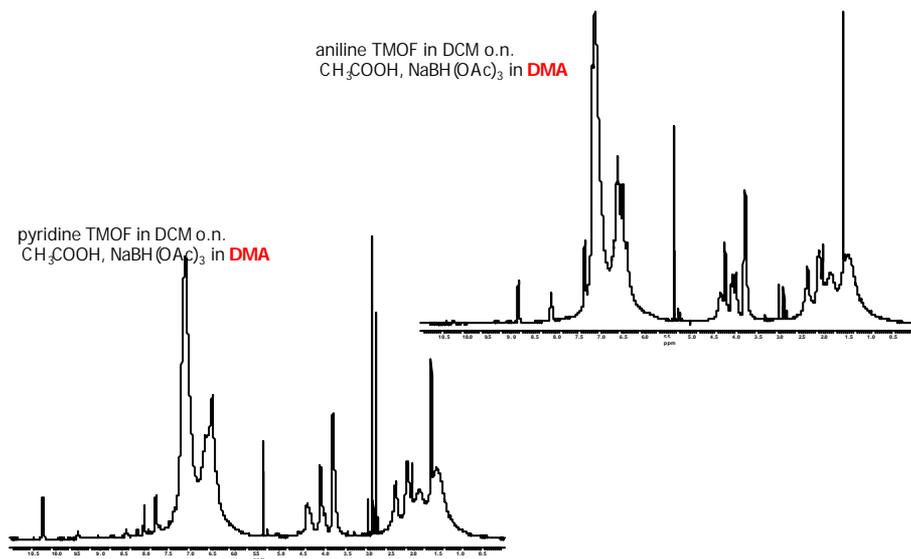
Pyridylamine: MAS NMR of a loaded resin



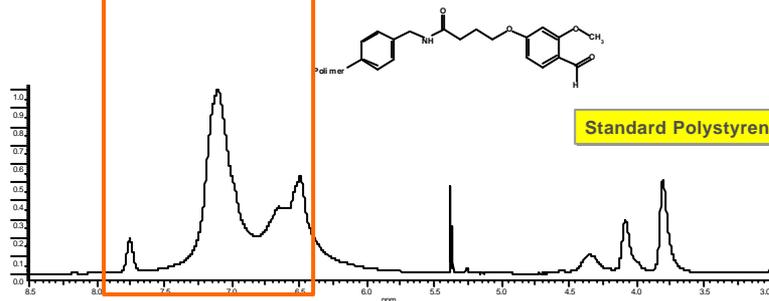
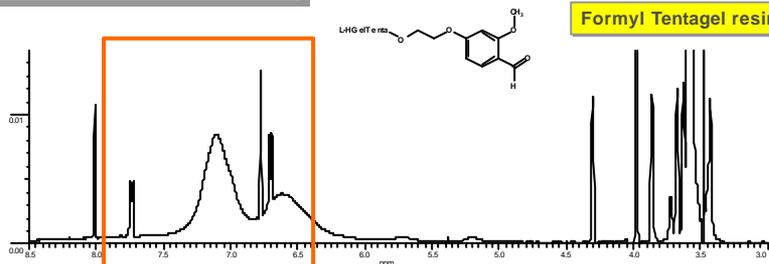
**Pyridylamine: MAS NMR of a loaded resin
Solid phase IR and HPLC of the cleaved product**



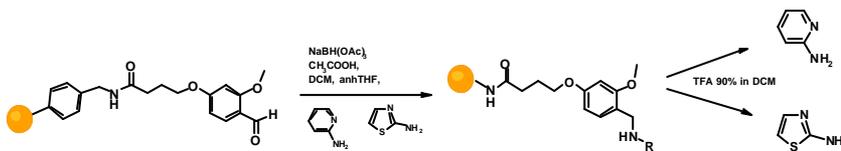
Some non obvious examples



High mobility solid support



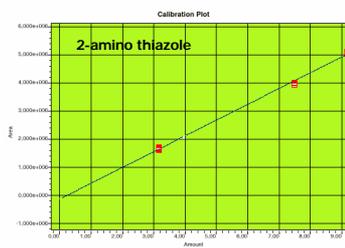
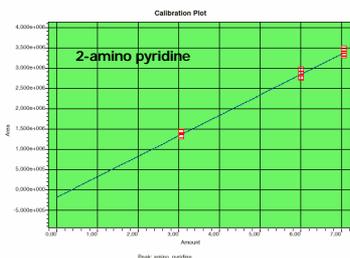
Yield calculation



We compared:

1. Weight
2. HPLC against standard,
3. Deconvoluted 1H MAS spectra

HPLC: calibration curve

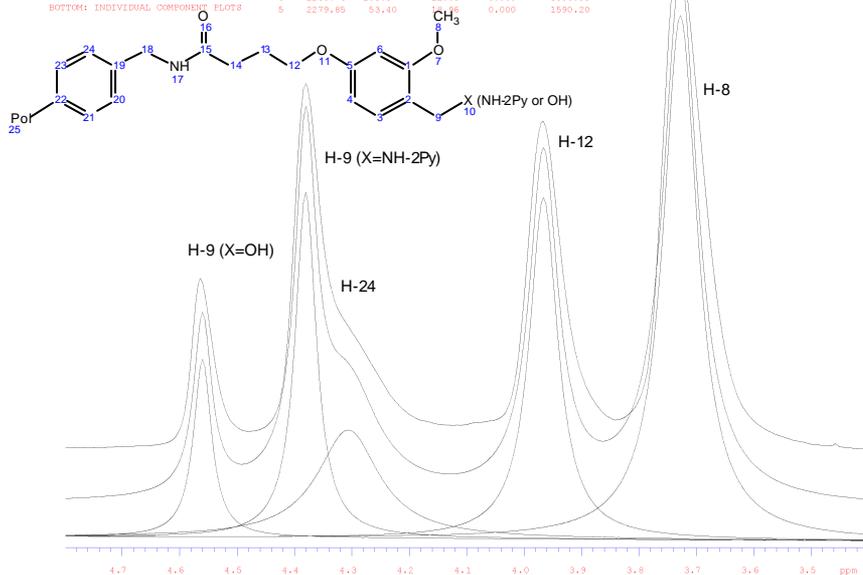


¹H-HRMAS-NMR: Deconvolution of Overlapping Signals

MULTICOMPONENT FIT, AUTOMATIC PLOT

LINE	FREQ (Hz)	HEIGHT	WIDTH (Hz)	GAUSS FR.	INTEGRAL
1	1862.49	157.07	36.27	0.000	8949.29
2	1982.74	102.29	30.05	0.000	4827.93
3	2151.41	32.42	65.26	0.000	3323.63
4	2188.74	103.57	22.05	0.000	3586.89
5	2279.85	53.40	44.25	0.000	1590.20

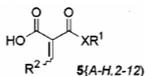
TOP: ACTUAL SPECTRUM
CENTER: FULL FIT
BOTTOM: INDIVIDUAL COMPONENT PLOTS



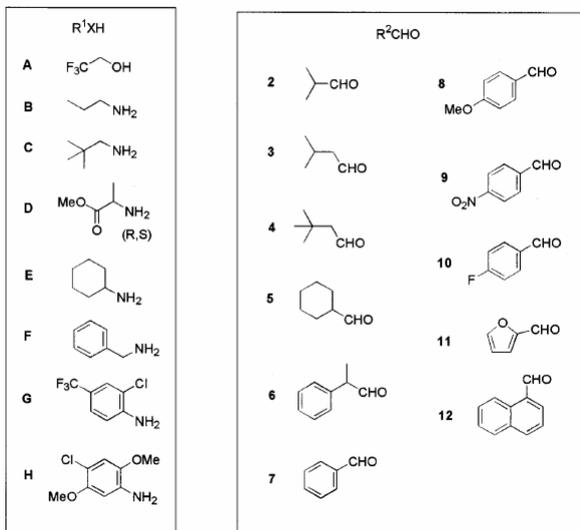
Yield evaluation by weight, HPLC against standard curve, deconvoluted NMR

entry	amine	resin mg (01.94m mol/g)	TMOF 10eq	CH ₂ COOH 1 eq	NaBH(OAc) ₃ 3eq	NaCNBH ₃ 3eq	solvent	NMR yield	WEIGHT Yield(mg)	HPLC Yield against Standard (%)
1	Thiaz	50		X	X		DCM	43%	5,9 (59%)	23
2	Thiaz	50	X			X	DCM	34%	10,6 (100%)	27
3	Thiaz	50	X			X	THF	n.d.	6,1 (60%)	2
4	Thiaz	50	X		X		DCM	40%	13,2 (131%)	32
5	Pyr	50		X	X		DCM	48%	11,8 (121%)	40
6	Pyr	50	X			X	DCM	53%	12,7 (123%)	43
7	Pyr	50	X			X	THF	35%	12,4 (121%)	27
8	Pyr	50	X		X		DCM	n.d.	6,2 (63%)	10

First implementations of rapid NMR quality assessments from titerplates



B. Hamper *et al.*

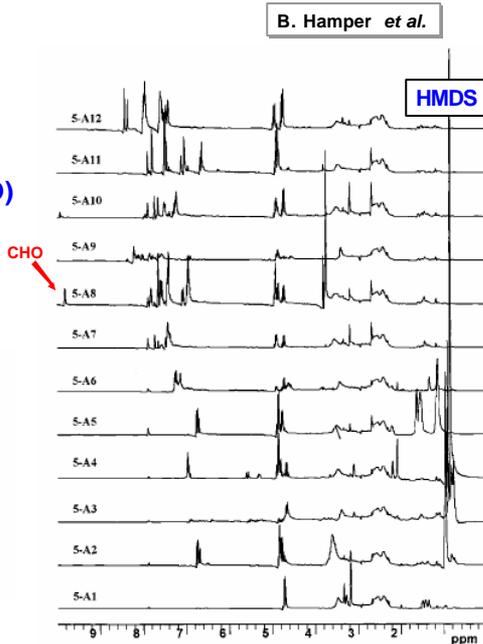
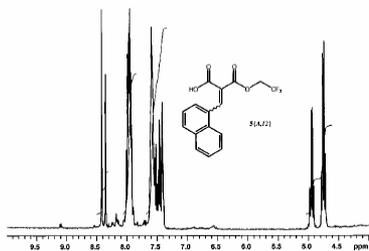


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Flow-through NMR Spectra in 96-well plate (regular DMSO)



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Complementary HPLC Analysis

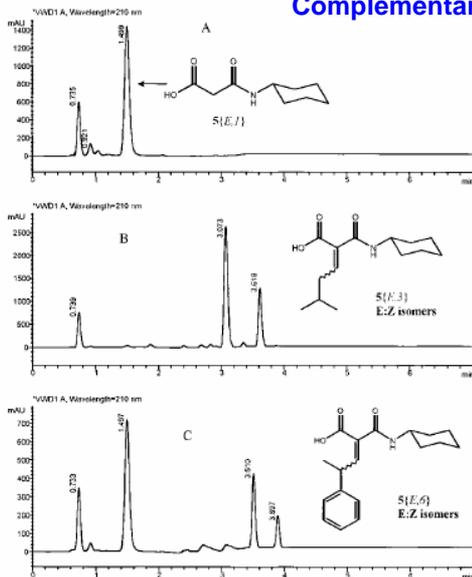


Plate View

Residual malonamide intermediates (%)

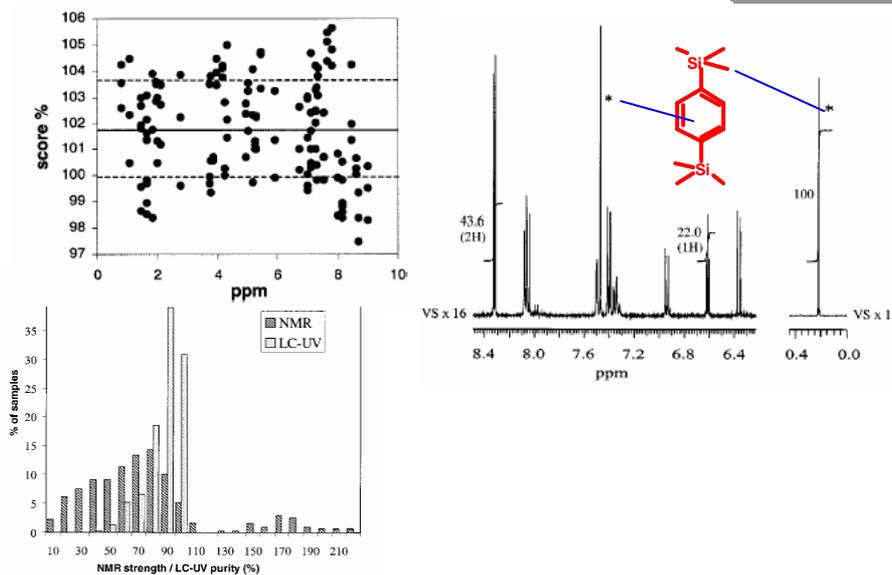
	1	2	3	4	5	6	7	8	9	10	11	12
A	100	0	0	0	0	0	10	5	0	15	5	0
B	100	0	0	0	0	100	0	0	0	0	0	0
C	100	19	0	0	7	94	6	5	0	6	0	0
D	100	0	0	0	0	90	0	0	0	0	0	0
E	100	7	3	0	4	48	0	0	0	2	2	2
F	100	14	11	4	12	100	11	8	3	3	5	5
G	100	13	8	5	16	24	8	10	5	8	0	5
H	100	35	22	22	20	100	18	16	12	21	0	22

Reference intermediates
Malonamides

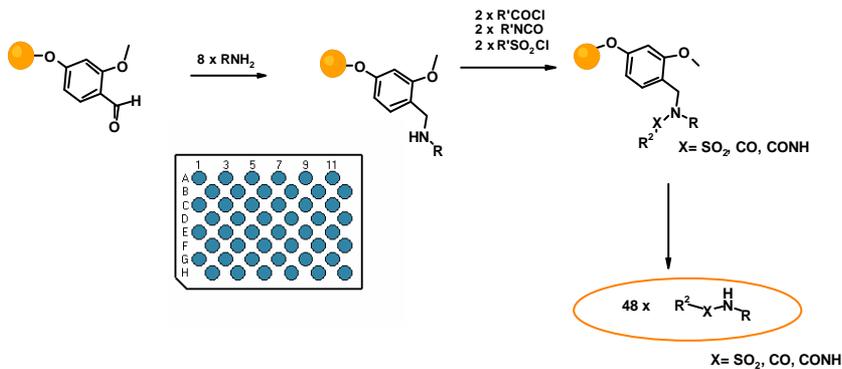
Development of accurate NMR quantitation with BTMSB standard

V. Pinciroli et al. *J. Comb. Chem.* 3, 434 (2001)

V. Pinciroli et al.



Application example (small compound array)



After cleavage the products were analysed by:

- HPLC-MS (identity, purity)
- ^1H NMR against standard (identity, purity, amount)
- ^1H NMR Vast System against standard (identity, purity, amount)



Direct-Injection NMR for High-Throughput Spectroscopy

VAST= Versatile Automatic Sample Transfer

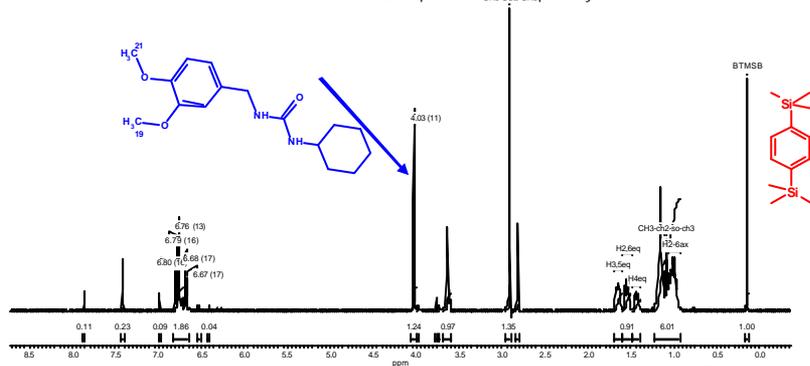
Experimental

4mM DMSO 400ml (H_2O presence does not matter)
 60 scans plus scout scan each sample
 20 min each sample (including temperature stabilization, gradient shimming on the first sample & 2 rinsing cycles per sample)

Virtually no contamination between samples

Reliable automation

2D expts can be run separately or in the run



NMR Routines on CombiChem Libraries

PHARMACIA

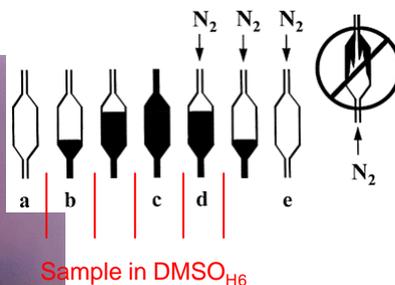
M. Tatò et al.

Direct-Injection NMR for High-Throughput Spectroscopy

96 wells plate run in nearly 16 hours

- 0.5-2 mg / 200 μl $\text{DMSO-}d_6$ (H_2O does not matter)
- 32-128 scans plus **scout scan** each sample
- 8 -12 min each sample (including temperature stabilization, gradient shimming on the first sample and 1 rinsing cycle per sample)
- virtually no contamination between samples
- reliable automation
- 2D expts can be run separately or in the run

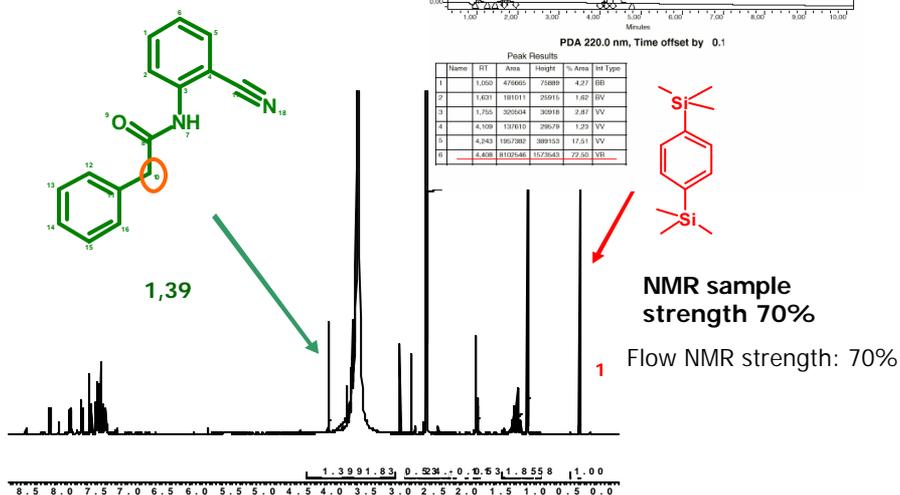
Direct-Injection NMR for High-Throughput Spectroscopy



VAST= Versatile Automatic Sample Transfer
(Varian)

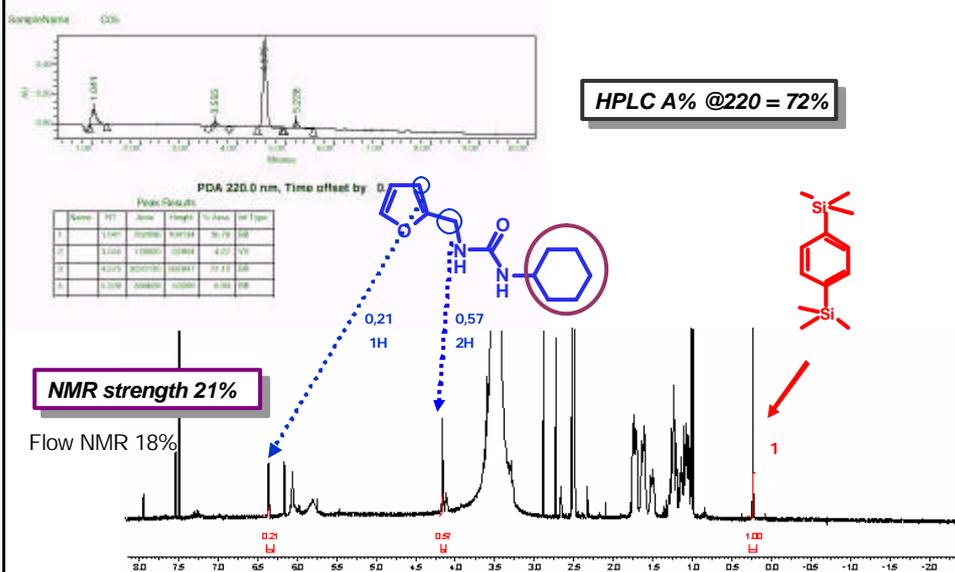
Ideal situation

HPLC A% @ 220 = 72%

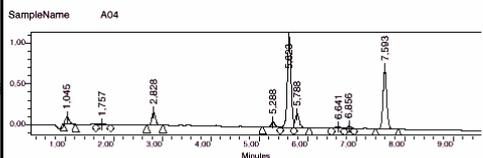


V. Pinciroli et al. *J. Comb. Chem.* 3, 434 (2001)

If the starting material is not UV visible



If there are no diagnostic NMR signals



HPLC A% @220= 46%

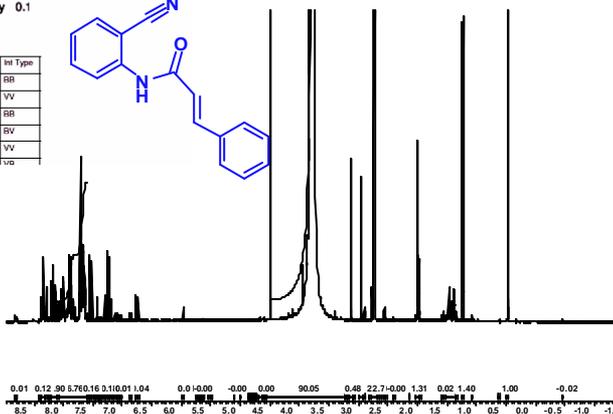
PDA 220.0 nm, Time offset by 0.1

Peak Results					
Name	RT	Area	Height	% Area	Int Type
1	1.045	4811021	88252	3.79	BB
2	1.757	120679	9637	0.36	VV
3	2.828	813403	160792	6.41	BB
4	5.288	353014	62733	2.78	BV
5	5.623	5842944	1095191	46.07	VV



NMR strength not easily measurable

Flow NMR not measurable



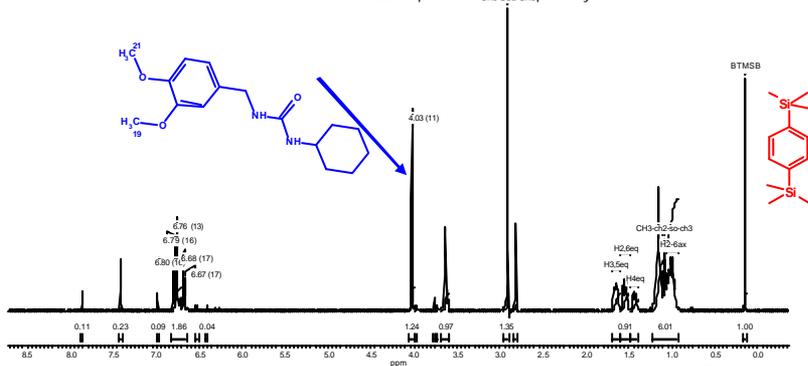
Direct-Injection NMR for High-Throughput Spectroscopy

VAST= Versatile Automatic Sample Transfer

Experimental

- 4mM DMSO 400ml (H_2O presence does not matter)
- 60 scans plus scout scan each sample
- 20 min each sample (including temperature stabilization, gradient shimming on the first sample & 2 rinsing cycles per sample)

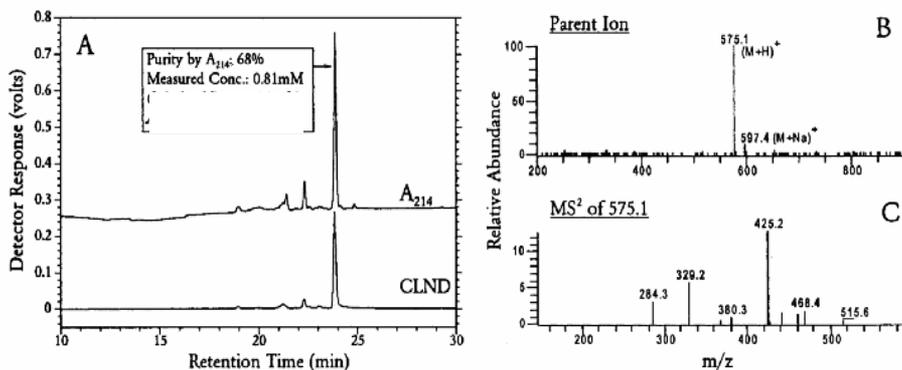
Virtually no contamination between samples
Reliable automation
2D exps can be run separately or in the run



Minimal Sample Consumption: Including Nitrogen Detection

HPLC/N/UV/MSⁿ detection platform

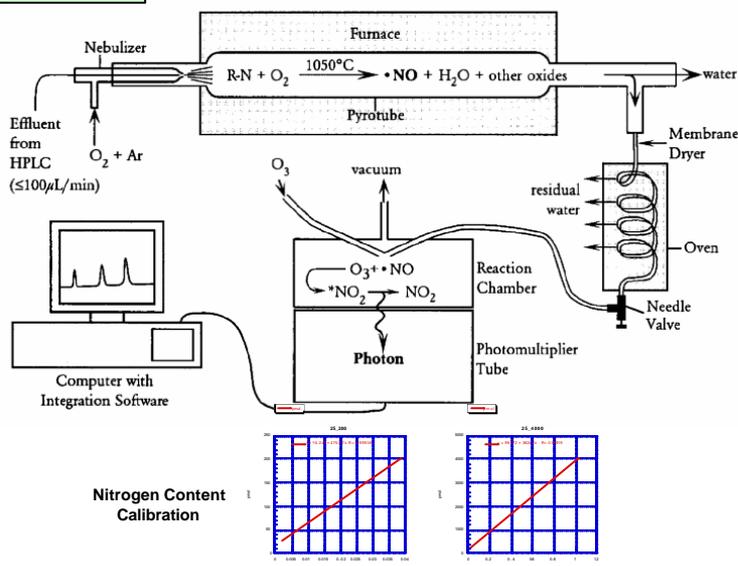
Identiv-Quantiv-Purity

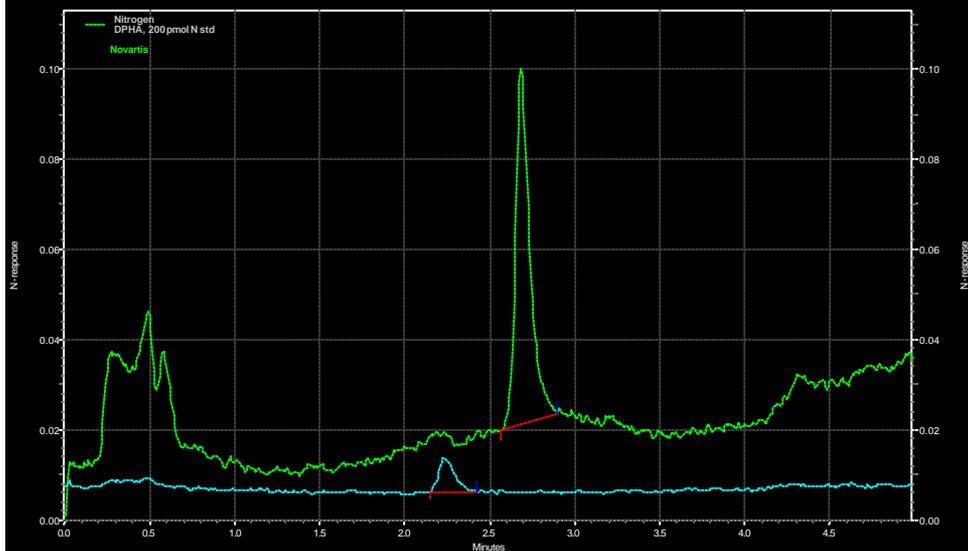


Integrated Analytics

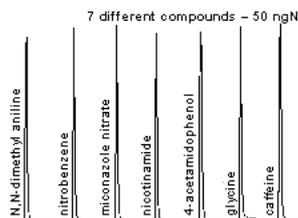
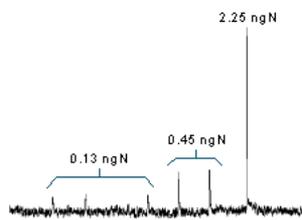
CHIRON TECHNOLOGIES
NOVARTIS

Nitrogen detection





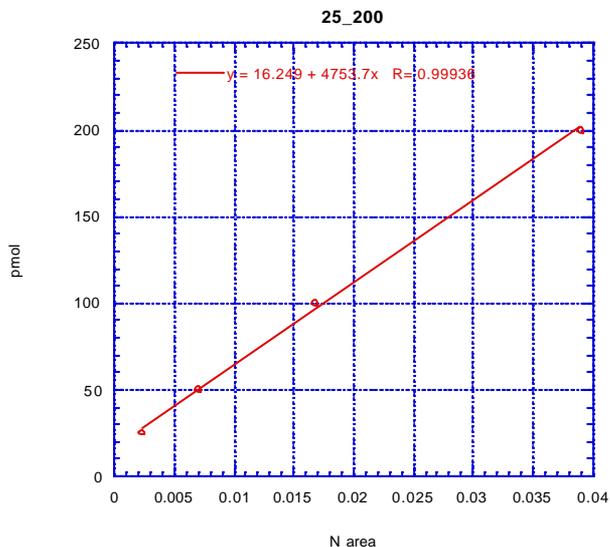
Nitrogen detection



Linear standard calibration curves over a $>10^4$ range of analyte concentrations.
 HPLC-CLND is compatible with reversed phase, size exclusion, and ion chromatography.
 Sensitivity is 0.5 ppm Gly-Gln peptide (0.1 ppm N) in water, $4x >$ absorbance at 215 nm.

Chemiluminescent reaction between ozone generated from oxygen and nitric oxide via high temperature pyrolysis of nitrogen-containing compounds

pmol Nitrogen Content



Concluding Remarks

- Derivatizing novel scaffolds on solid phase with standard reactions is an efficient, compound prolific approach
- Ample choice of techniques for in-process control of solid phase syntheses (on and off beads)
- No significant bottlenecks
- Beware quantitation by weighing of non-crystalline microsamples (< 5mg)
- NMR quantitation against standard is the most accurate system however ... it is relatively slow. Impractical for samples < 1mg
- The MAS NMR on solid phase gives good qualitative information, the quantitative evaluations may be less precise and more demanding in terms of interpretation (spectra deconvolution)
- Nitrogen detection (HPLC online) has excellent throughput and is the method of choice for the quantitation of microsamples of large libraries
- Initial concerns on analytical limitations of solid phase chemistry have been overcome

Acknowledgements

Katia Martina - *Combinatorial Chemistry*

Marco Tatò - *Structural Chemistry*

Roberto Biancardi - *Structural & Predevelopment Analysis*

Bruce Hamper - *Combinatorial Chemistry, St. Louis USA*

The Chemistry Department – Pharmacia Italy, Nerviano (Milano)