Molecularly Imprinted Polymers

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2010年11月30日(東京)と12月2日(大阪)に開催した ExploraSepセミナーのプレゼンテーション資料です。



Uppsala: Summer and Winter

















バイオタージ本社があるスウェーデンの風景

Agenda

Part I

- ➤ What are MIPs a short history?
- ➤ How do they behave?
- > Examples of selectivity
- ➤ ExploraSep™: a new screening concept

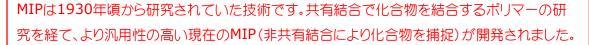
Part II

➤ ExploraSep and Genotoxins: Case Studies



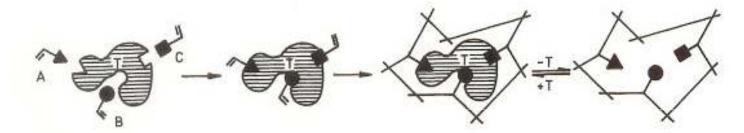
The concept of 'Imprinting'

- In 1931 the group of Polyakov (Kiev) reported some unusual adsorption properties in **silica particles** prepared using a novel synthesis procedure. Sodium silicate had been polymerized in water using (NH₄)₂CO₃ as the gelating agent. After two weeks, **additives (benzene, toluene or xylene)** had been added. The silica was subsequently allowed to dry for 20–30 days, after which the **additive was removed** by extensive washing in hot water. Subsequent **adsorption studies** revealed a higher capacity for uptake of the additive by the silica than for structurally related ligands, i.e. some kind of **memory for the additive was apparent**, at least in the cases of benzene and toluene.
- In 1934-5 more detailed investigations of the phenomenon were reported and the observed selectivity was explained as resulting from structural changes in the silica reflecting the nature of the additive.
- 1950,s Polyakov repeated and reiterated his founding experiments though they were not widely accepted or referenced
- 1970's Gunther Wolff (Germany) took the concept to another level....

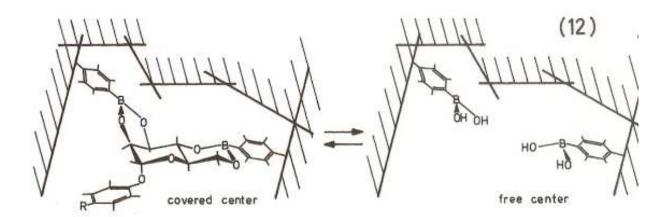


The Birth of Covalent Imprinting Wolff (1972-78)

"Polymers with binding groups located in a definite spatial proximity and cooperativity in cavities of specific shape should show high selectivity in binding. This arrangement is similar to those of natural receptor sites."



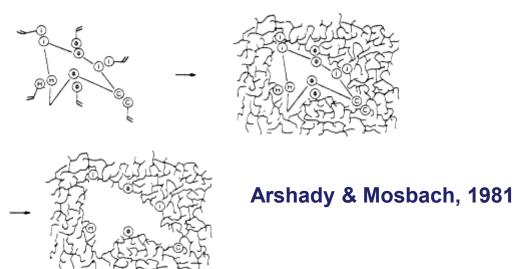
Covalent imprinting – boronate esters with diol templates





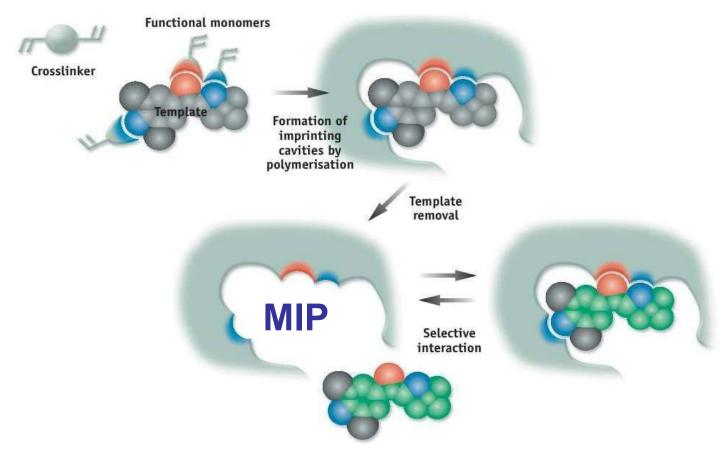
Non- covalent Imprinting – the universal method

"According to this strategy a monomer mixture containing ...cross-linking units is polymerized in the presence of a free substrate ..act as a template.... This is simply a mixing process and no chemical attachment to the monomeric units is required. The monomers are, however, chosen in such a way as to have non-covalent binding abilities (ie ionic, hydrogen bond, hydrophobic, charge transfer etc) complementary to those of the guest template."



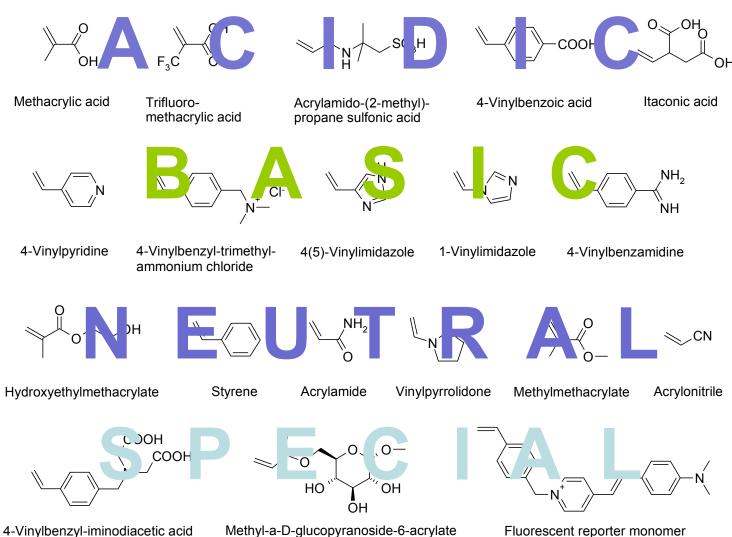


The Non-covalent Process simplified

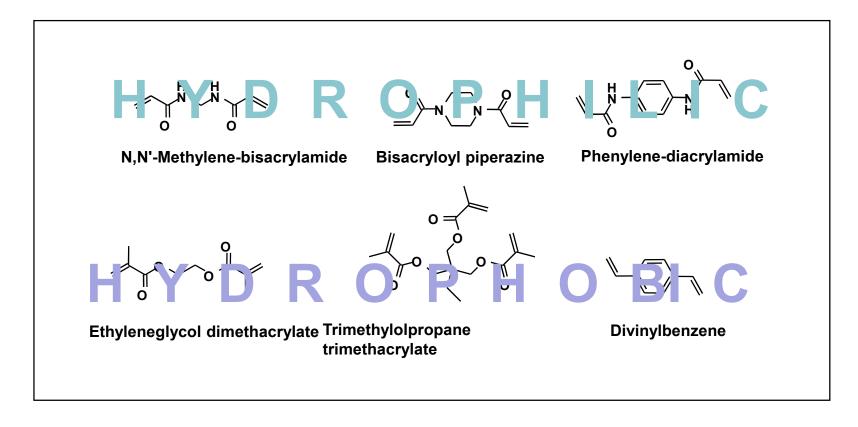




The Functional Monomer Landscape



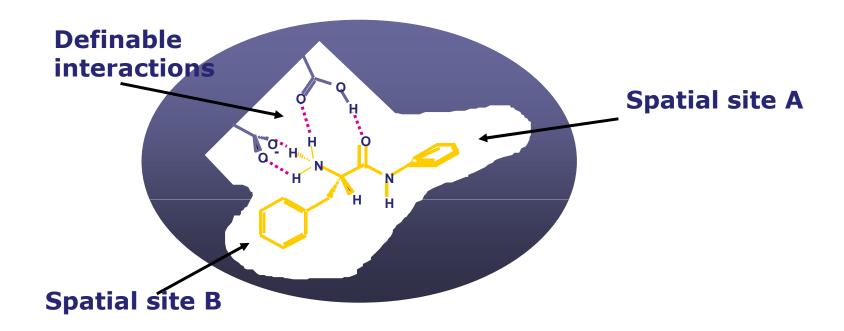
The Polymer backbone







The MIP Binding Site (small molecule)



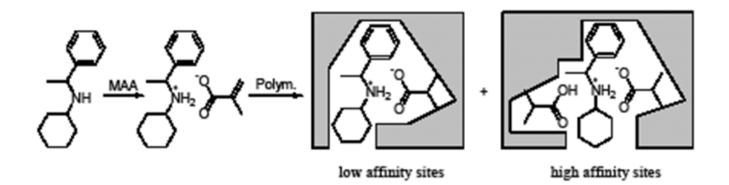
With the monomers employed interactions can involve H-bonding, charge-charge, van der Waals, hydrophobic and charge-transfer

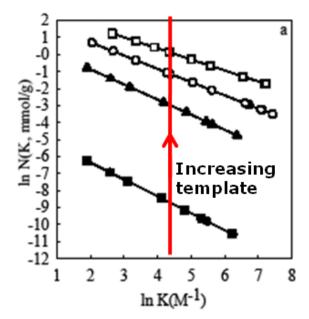


Engineering binding and Capacity



Template ratios: K and Capacity



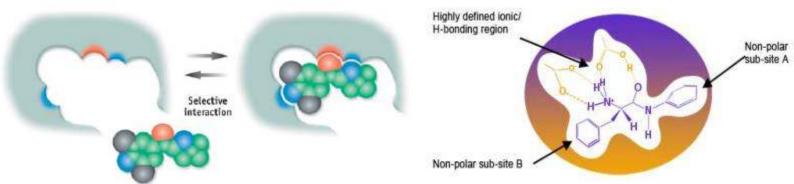


- 1) Template-monomer stoichiometry controls capacity
- 2) The random polymerization process generates polyclonal sites



Molecularly Imprinted Polymers – a Summary

- Highly cross-linked polymer based phases
- Pre-determined selectivity for a particular analyte or group of structurally related compounds
 - Size exclusion
 - Chemical interactions in highly defined positions (H-bonding/ionic, Van der Waals interactions, π – π interactions)
- Selective target recognition
 - Mimics of <u>polyclonal</u> antibodies or receptors





MIPは高度に架橋された特殊なポリマーで、特定の化合物をかたどったキャビティ内においてポジション限定的に化合物と相互作用します。選択的に化合物を"認識"する仕組みは抗原抗体反応(ポリクローナル)に似ています。

Agenda

Part I

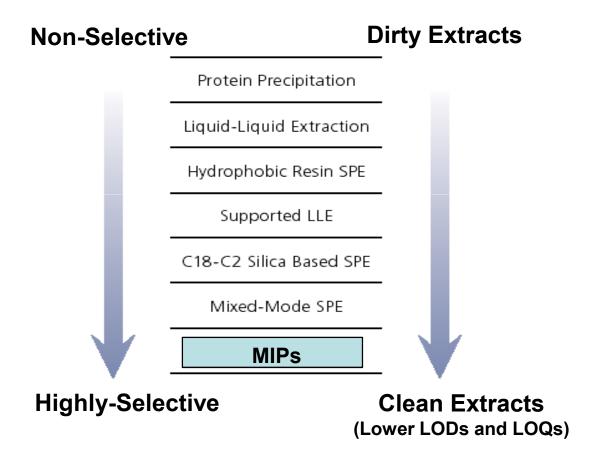
- > What are MIPs?
- ➤ How do they behave?
- > Examples of selectivity
- ➤ ExploraSep™: a new screening concept

Part II

> ExploraSep and Genotoxins: Case Studies

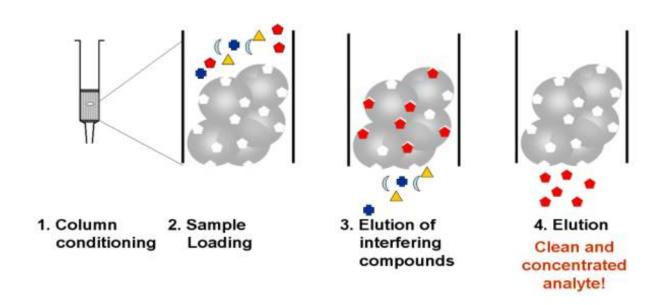


Relative Selectivity of Sample Preparation Techniques





AFFINILUTE MIP SPE procedure



- MIP methodology differs from conventional SPE methodology
- Protocols for reversed phase, ionic-exchange resins etc CANNOT be used without careful comparison with the recommended MIP method
- Selectivity is typically introduced during the interference wash step with organic solvents



AFFINILUTE MIP: Selective extraction in complex matrices

Technical features

- MIPs generate stronger interaction between the sorbent and the analyte
- Interfering substances can be washed away using harsher washing conditions
- The MIP material is stable at high temperatures, in organic solvents and at extreme pH (normally)

Advantages

- Cleaner extracts
- Simplified washing protocols with less extraction steps
- Minimized matrix effects and ion suppression
- Higher precision



AFFINILUTE MIP Phases and Applications Commercially Available For...

Drug-like

- Clenbuterol
- Beta agonists (class-selective)
- Beta blockers (class-selective)
- Beta receptors (class-selective, β-agonists & β-blockers)
- NSAIDs

Forensic

- Amphetamines
- NNAL
- TSNAs (Tobacco Specific Nitrosamines)

Food/Agricultural

- Chloramphenicol
- Fluoroquinolones
- Nitroimidazoles
- Triazine herbicides (class-selective)
- Polyaromatic hydrocarbons (PAH)



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Example 1: Amphetamine drug Family

- A class of compounds including amphetamine (alpha-methylphenethylamine) and substituted amphetamines
- CNS stimulants
 - Clinically used to treat ADHD, narcolepsy and other sleeping disorders.
 - Stimulants and hallucinogens illegally used as recreational club drugs and as performance enhancers.
- Heavily regulated worldwide
- Schedule I & II drugs as reported by the DEA, USA



Amphetamine Structures

Methamphetamine "crystal meth", "ice" – potent stimulant MDA, MDEA & MDMA (Ecstasy) – psychedelic stimulant

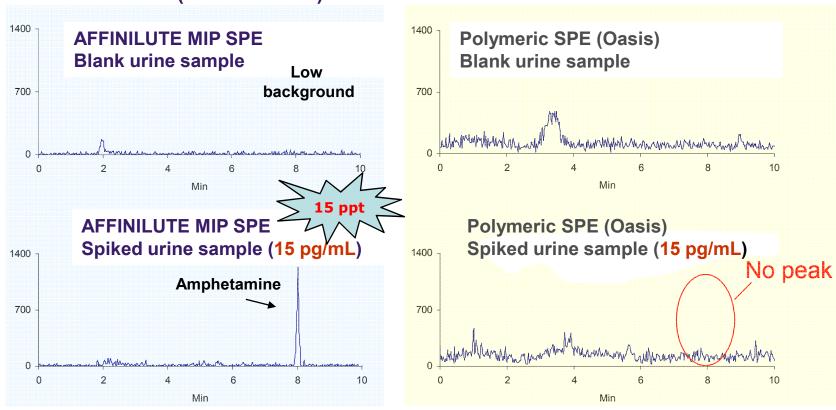
MDA (3,4-Methylenedioxyamphetamine), MDMA (3,4-methylenedioxy-N-methamphetamine) MDEA (3,4-methylenedioxy-N-ethylamphetamine)



AFFINILUTE MIP: High Sensitivity

AFFINILUTE MIP vs Conventional Hydrophilic Polymer[†]

Urine extracts (MRM 136/118)



Higher recovery and reproducibility

AFFINILUTE MIP vs Conventional Hydrophilic Polymer[†]

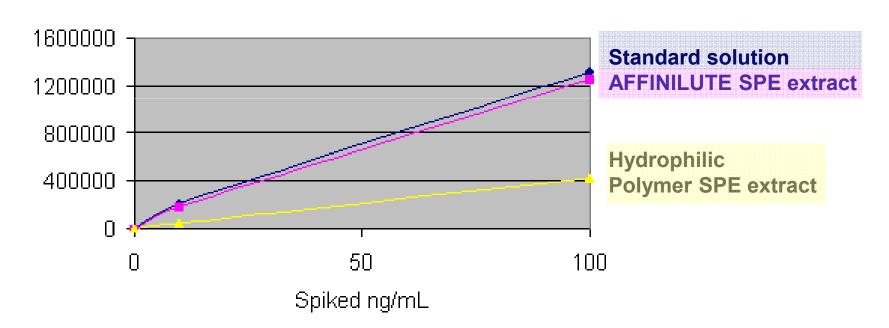
	% Recovery		% RSD	
	Affinilute	Hydrophilic polymer SPE	Affinilute	Hydrophilic polymer SPE
Methamphetamine	101	100	1,41	5,16
Amphetamine	104	90	3,90	14,30
Phentermine	104	64	6,11	26,36
MDA	113	÷	9,84	*
MDMA	97	86	2,52	8,10
MDEA	106	7	6,60	37,80
	High Recoveries for all compounds using AFFINILUTE		High Reproducibilit using AFFINILUTE	у



Lowered Ion-Suppression

AFFINILUTE MIP vs Conventional Hydrophilic Polymer

Ion suppression, MDEA





Example 2: NNAL

- Carcinogenic nicotine metabolite measured in urine of smokers and passive smokers
- A nitrosamine so within the genotoxic 'class'
- Current methods elaborate and timeconsuming
- Trace level determination required











NNAL Structure

Note: -N-N=O is a PGI (genotoxin) functional group



Affinilute™ NNAL

Conventional method

Analysis of NNAL. To 20-100 ml (smokers) or 100-500 ml (non-smokers) of urine, 4 ng iso-NNAL (internal standard, generous gift from Dr Dietrich Hoffmann, Valhalla, NY, USA) were added. Samples from non-smoker were concentrated on a rotary evaporator at 40 °C and 3 kPa to 100 ml. The sample was adjusted to pH 5.0 with hydrochloric acid. The aqueous solution was transferred to a glass column (450 mm × 40 mm, G3 frit) filled with Extrelut® (Merck, Darmstadt, Germany) and allowed to soak for 30 min. The column was eluted with 250 ml ethyl acetate (Code 3427, Promochem, Wesel, Germany), and the eluate was evaporated to 5 ml at 40 °C and 20 kPa. The concentrate was added to 5 ml water, adjusted to pH 2.0 with hydrochloric acid, and transfered to a separation funnel. After washing the aqueous layer three times with ethyl acetate, the pH was adjusted to 5.0 with aqueous sodium hydroxide and absorbed on a column containing Extrelut®. The column was eluted with 80 ml ethyl acetate, the eluate dried over anhydrous sodium sulphate, and concentrated to 2 ml in vacuo. The final extract was applied to 8 g aluminium oxide (activity II-III, ICN Biomedicals, Eschwege, Germany), equilibrated with 20 ml ethyl acetate (glass column, 150 mm × 15 mm, G2 frit). The column was washed with 10 ml ethyl acetate. Unconjugated NNAL was eluted with 20 ml ethyl acetate/methanol (10:1 v/v) (methanol, Code 9835, Promochem, Wesel, Germany) and evaporated to 1 ml. The purified fraction was transfered to a reaction vial, the solvent was removed under nitrogen, and the residue was derivatized by adding 48 μl bis-(trimethylsilyl)acetamide (BSA) and 2 μl trimethylchloro silane (TMCS) (Aldrich, Steinheim, Germany) at 50 °C for 30 min.

Over 20 laborious steps
Results in 3 days
Insufficient detection limits

Affinilute method

Extraction Procedure:

Recommended flow rate is 0.5 mL/min except for analyte elution 0.2 mL/min. Gentle vacuum should be applied between each interference elution.

Sample pre-treatment: Urine diluted 1:1 with dest. water

Column conditioning: Condition the column with:

1 mL of DCM

1 mL of MeOH
 1 mL of dest, water

Sample application: Apply the sample to the column (1-10 mL)

Interference elution: Elute the interferences with:

 1 mL of water (elution of salt and matrix components)

 10 minutes of vacuum (~ -0.7 bar) to dry the column

1 mL of toluene

1 mL of toluene/DCM (9:1)

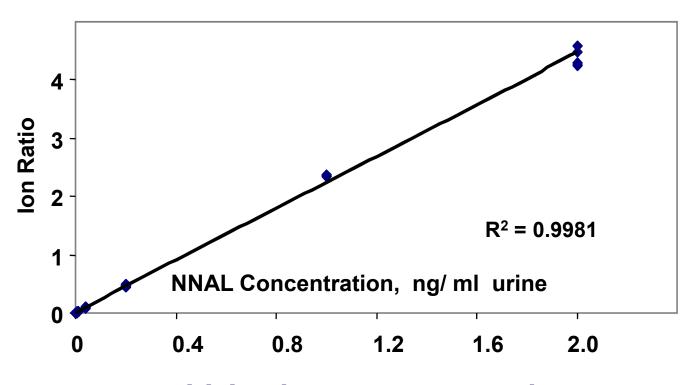
 1 mL of toluene/DCM (4:1) (selective wash, elution of hydrophobic bonded interferences)

2 minutes of vacuum to remove the toluene

Results after 0.5 h!



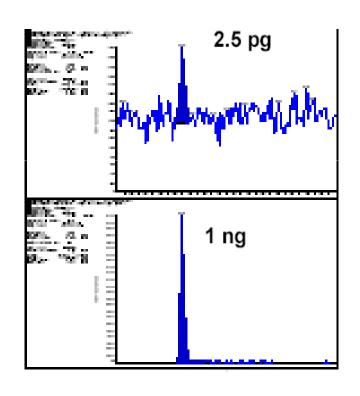
Affinilute™ NNAL - sensitivity



Sensitivity down to 0.1-0.2 ppb



Affinilute™ NNAL – the sensitivity LIMI





John Bernert, CDC Atlanta

Faster and more efficient SPE by using MIP allows higher sample throughput

Allows high sensitivity – detection of 2 ppt possible!



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ExploraSep[™]: Exploiting Similarity

ExploraSep ► more...

Screening Plates

Highly specific scavengers for removal of impurities including genotoxins



Target compounds are screened on multiwell plates containing a large number of different MIPs generated against many different templates and containing different monomer chemistries

How and why does ExploraSep work?



Similarity – the Concept

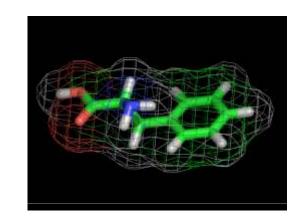
- Virtually all ways to measure molecular similarity are based on either (or both!) of these things:
 - The structure of the molecule (2D, 3D)
 - The properties of the molecule
- In general, an object A (molecule) is described by a set of n features:

$$X_A = \{X_{1A}, X_{2A}, X_{3A}, ... X_{nA}\}$$

This is generally known as a fingerprint



- A similarity measure or
- A dissimilarity (distance) measure



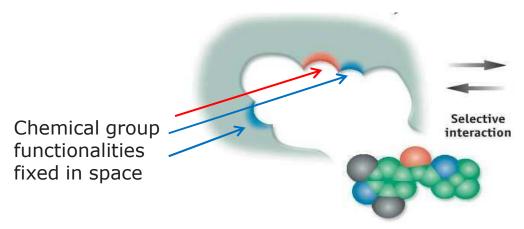
Most of the time we use the concept of similarity because of the similar property principle:



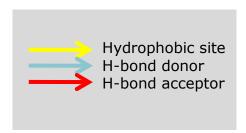
Structurally similar molecules are expected to exhibit similar physical properties or biological activities

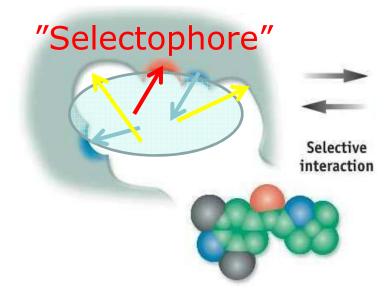


Similarity leads to cross-reactivity



Other molecules that share common features may also fit into the MIP binding site







Selectophore concept Summary

- Comparable to <u>pharmacophore</u> concept in medicinal chemistry*
- Pharmacophore ="a set of structural/chemical features in a molecule that is recognized at a receptor site and is responsible for the biological activity"
- Similar molecules can bind to the same site otherwise Pharma would have no business!
- It is to be expected that the selectivity 'profile' of a MIP binding site will be at least as flexible as a biological receptor site

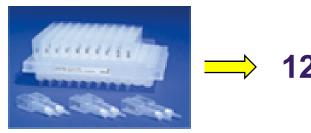
*Gund, P., *Prog. Mol. Subcell. Biol.* **1977**, 5: pp 117–143



ExploraSep Chemistries

ExploraSep ► more... Screening Plates

Highly specific scavengers for removal of impurities including genotoxins



⇒ 128 resins

Plate A (acidic functionalities)

- Hydrophobic and carboxylic acidic moieties
- Successful target analytes amines, amides, nitrosamines, esters, carboxylic acids

Plate U (proprietary 'urea' functionalities)

- Hydrophobic moieties and urea groups
- Successful target analytes phosphates, phosphonates, sulphates, sulphonates, anions of carboxylic acids

Plate H (aromatic functionality)

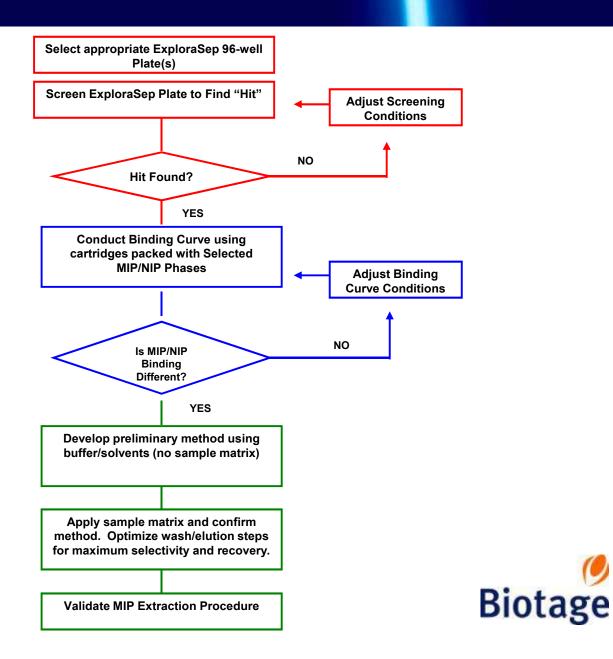
- Hydrophobic aromatic moieties
- Successful target analytes non-polar and aromatic compounds

Plate C (CHO or multi-OH binding functionality)

- Hydrophobic moieties and hydroxy functionalities
- Successful target analytes 1,2- and 1,3-diols, and α-hydroxy carboxylic acids

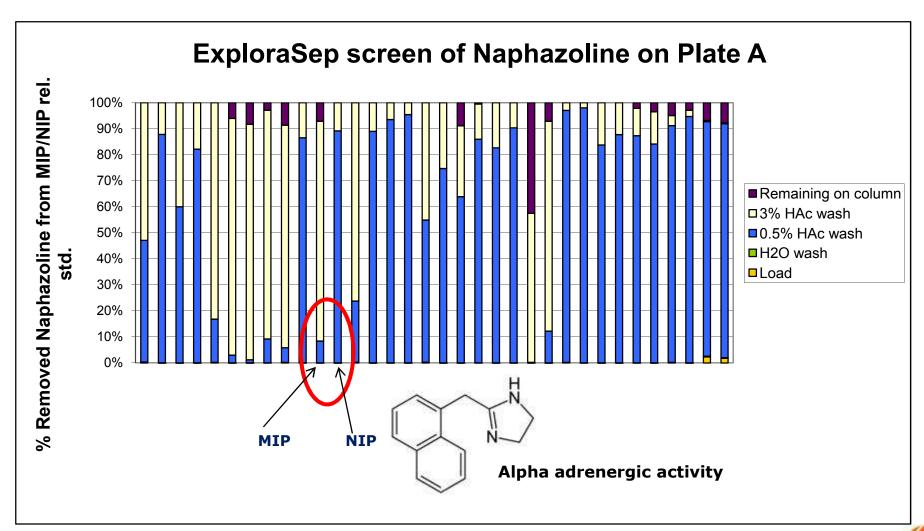


Overview of ExploraSep Process



ExploraSepによるMIPスクリーニングから、 最終的に実用のためのプロトコルを決定する までのプロセスの概要です。

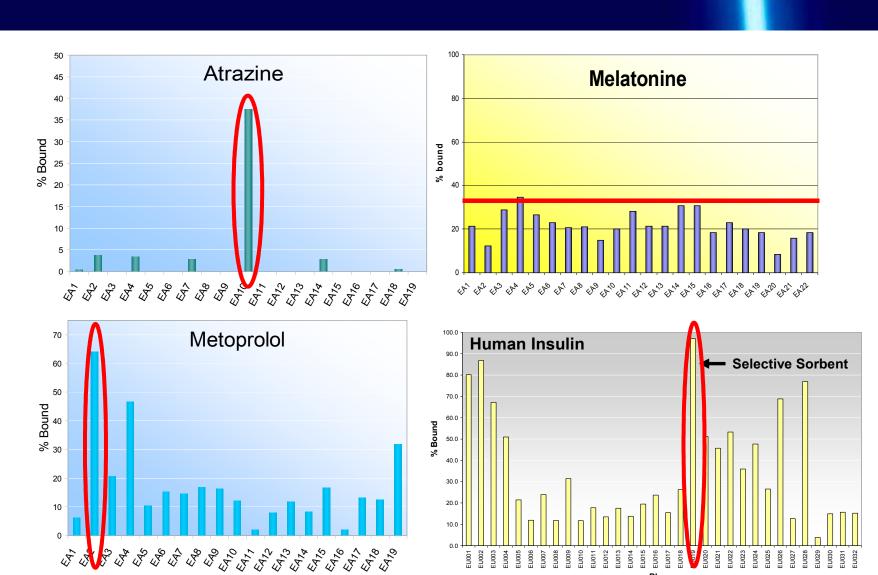
ExploraSep Screening: The Process



ExploraSepには、各種MIPと、それぞれのMIPに対応するNIP(non-imprinted polymer)が載っています。NIPは、MIPと全く同じモノマー組成で、テンプレート(鋳型)化合物を使わずに作成したポリマーです。



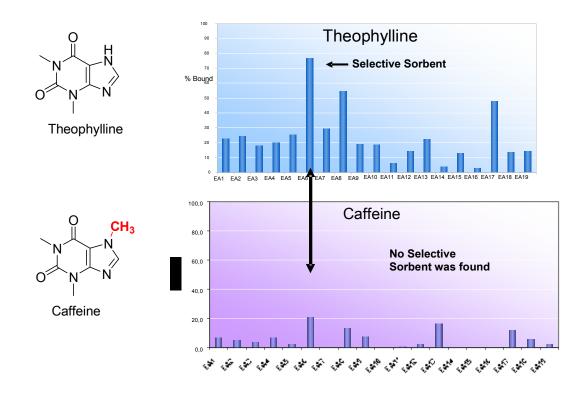
ExploraSep Screening Examples



Biotage



ExploraSep: Sensitive to small differences



A MIP that binds theophylline but not caffeine Differential binding with only one methyl group difference!



Summary

- MIPs are artificial binding sites in polymers and can be made to operate in SPE or chromatography mode
- MIPs are **stable** to all organic solvents and normally extremes of pH (depends on monomer chemistry)
- MIPs can be made selective for a restricted group of highly related molecules OR a 'CLASS' of chemically similar compounds
- MIPs can be manufactured for use at process scale
- New separation materials for most compounds of interest to pharma can be discovered using ExploraSep



End of Part I



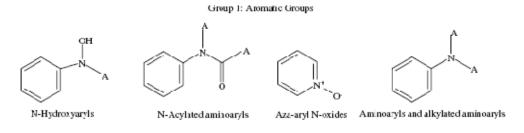
Agenda

Part I

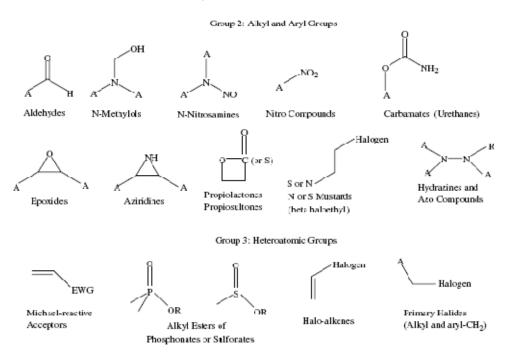
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Potential Genotoxic Impurities (PGI): Structural Alerts



Purines or Pyrimidines, Intercalators, FNAs or PNAHs



(From Müller et al, Regulatory Toxicology and Pharmacology 44 (2006) 198–211)



Genotoxic impurities and ExploraSep™

The Analytical Challenge

- Validated methods for multiple API's and PGI are required
- The genotoxic compounds may be closely related to the API

The Process Chemistry challenge

- Investigate process conditions that may create PGI
- Map the effect of all reaction conditions that effect PGI formation
- Investigate process changes required to control PGI concentrations
- Develop back-up strategy to scavenge PGI with high yield of API



PGI Screening Process

ExploraSep

- Screen each of 4 plates in clean solvent mixture containing API and PGI
- Take 'hits' and develop separation method (SPE or chromatography

Comprehensive data

 Raw data supplied with analysis showing polymer of choice for further development

Rapid Results

Projects completed on agreed schedule at fixed price

Proven Scale-up

 MIP production already at 500 Kg/year. Option of bulk resin, SPE cartridges, pre-packed Flash cartridges or HPLC preparative columns

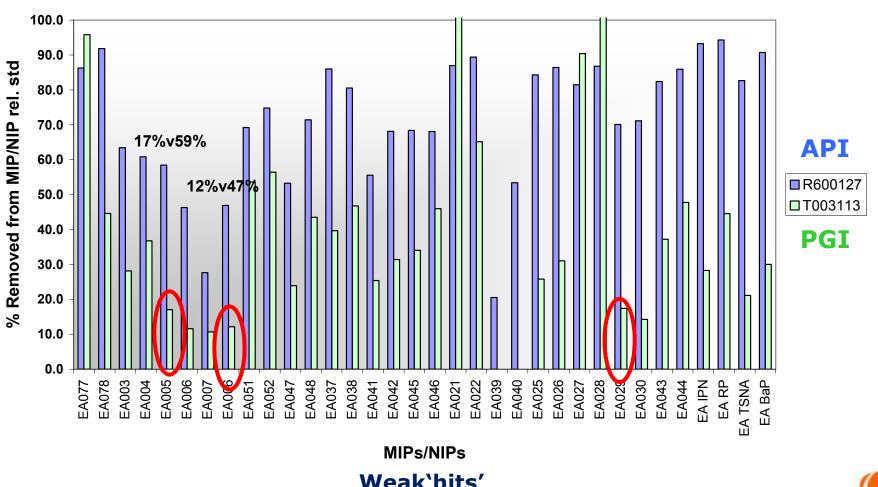


Case Study 1 A large US Pharma



CS 1: Screening for Genotoxin over API

ExploraSep Plate A: loaded in toluene

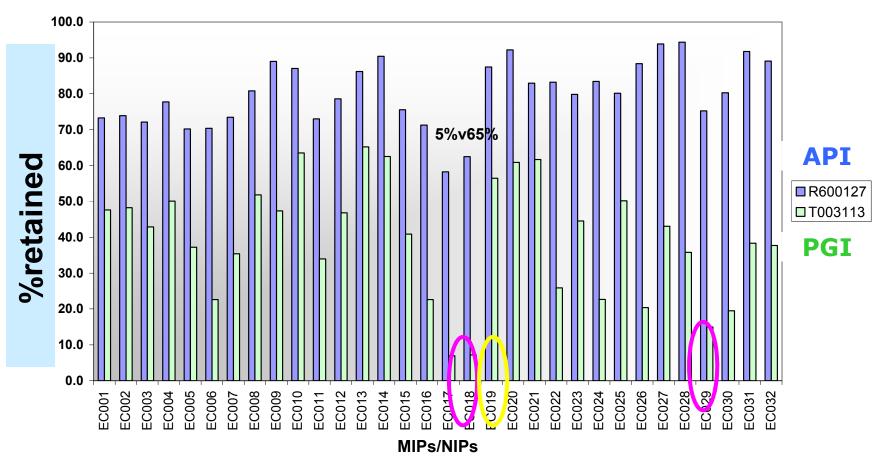


Weak'hits'



CS 1: Selectivity for Genotoxin over API

ExploraSep Plate C: loaded in toluene

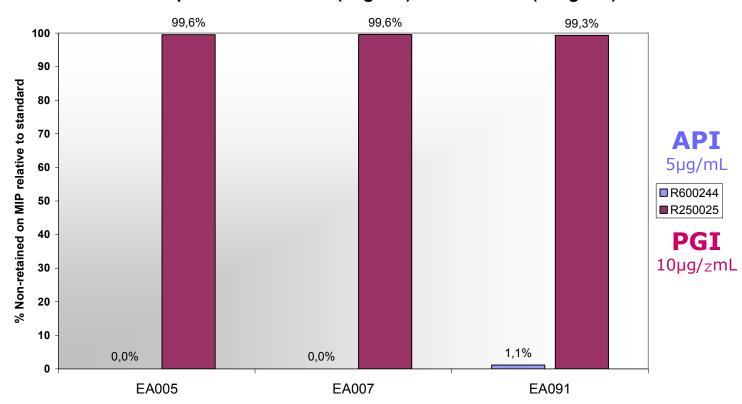


Several hits that differentiates PGI (T003113) and API (R600127)



CS 1: API binds and PGI elutes

Elution profile of R600244 (5ug/mL) and R250025 (10ug/mL)



High Selectivity forR600244 (API) over R250025 (PGI)

(0% = non-detectable by lc/ms/ms)



CS 1: Conclusion

- First API-PGI set screened on Plate C identifies several candidates with selectivity for PGI for further evaluation. EC 029 selected for further evaluation
- Second API-PGI set screened on plate A shows selectivity for API over PGI. EA 005 and EA 007 selected for further evaluation.



Case Study 2 A large European Pharma

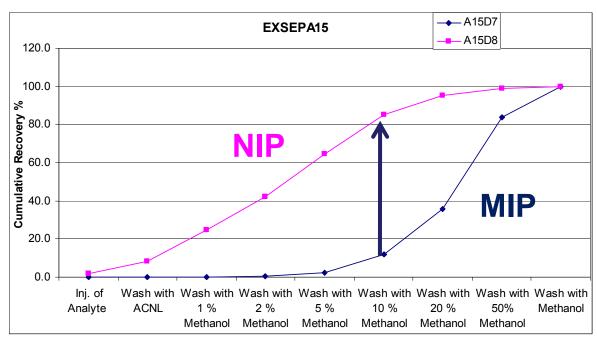


Case Study 2

- Structures confidential but:
 - PGI contains amide and piperazine moieties.
 - API contains piperazine, indole and dione moieties.
- Log P
 - PGI ~ 1.3
 - $API \sim 3.3.$



Case Study 2



Binding curve for genotoxin (d7) and API (D8) on MIP EA15 derived from ExploraSep A Plate.

Arrow shows difference between imprinted (MIP) and non-imprinted (NIP) polymer



Conclusion

- Quite good selectivity on several of the polymers on the A plate: EA07, EA15 and EA23
- Polymers were packed in HPLC columns and supplied to customer



Case Study 3 A European CMO



Case Study 3

- All genotoxin structures known but only a few of the APIs.
- Log P of Methyl p-Toluenesulfonate (PGI) ~ 2.0, log P of 21-chlorodiflorasone (API) ~ 2.7.
- Log P of 1,3-diisopropylurea (PGI) ~ 0.6, log P of API ~ 2.7.
- All screened with ExploraSep



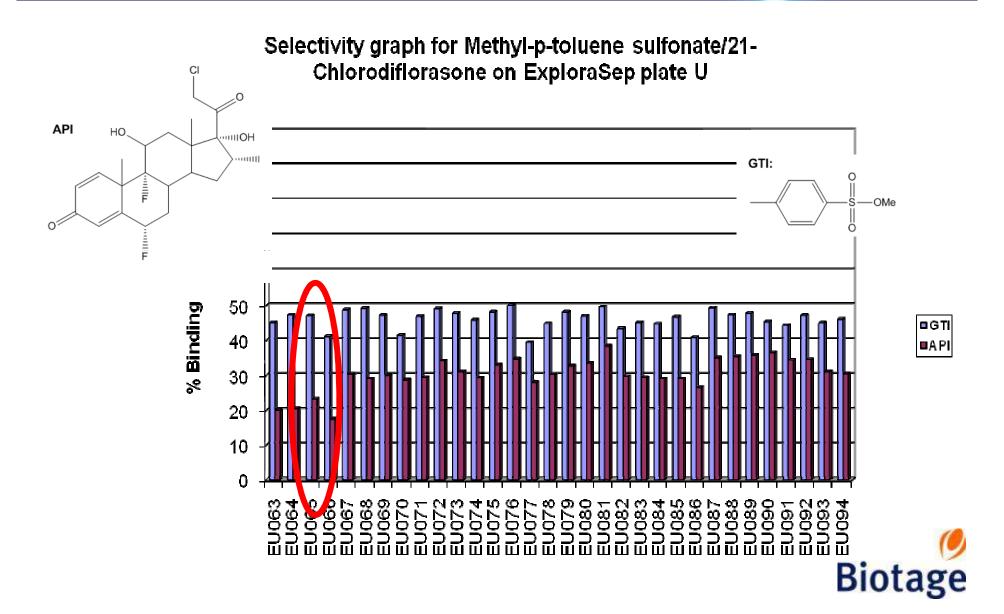
API or Intermediate:

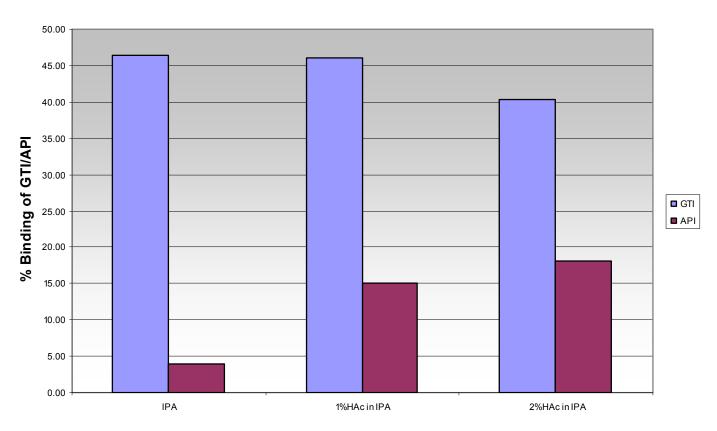
Methyl p-Toluenesulfonate

GTI:

1,3-diisopropylurea



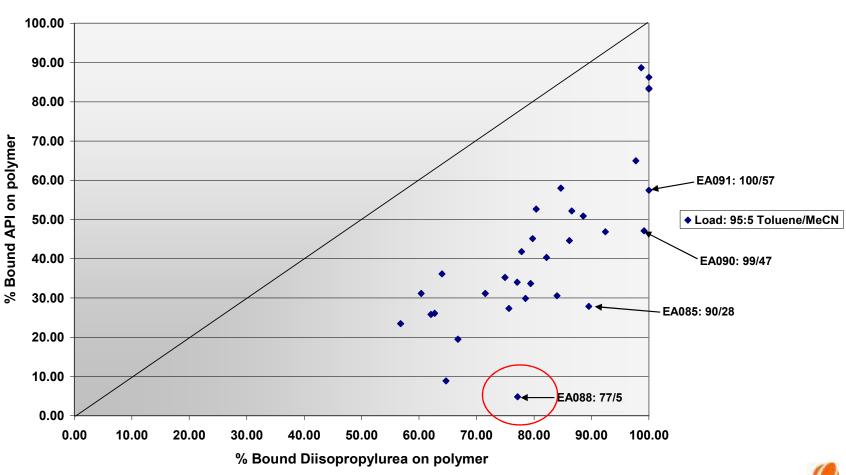




Selectivity graph of methyl p-toluenesulfonate (PGI)/API in loading on EU064



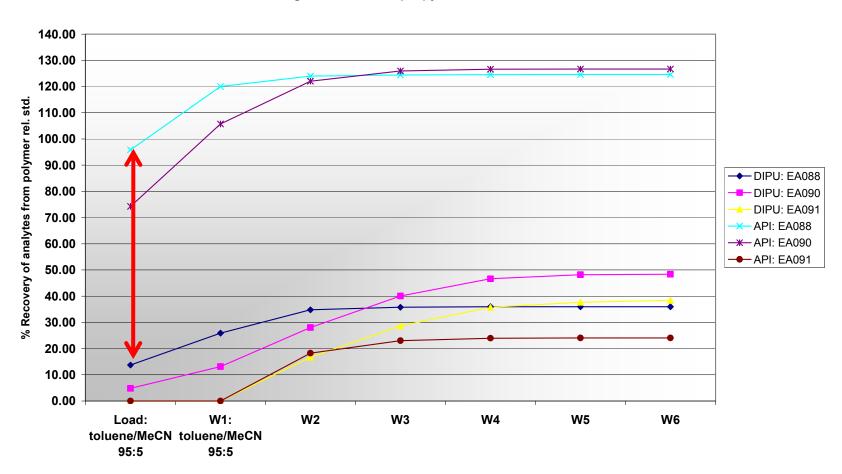
Selectivity Plot of Diisopropylurea vs. API on ExploraSep plate A in Loading



Further evaluation of additional A plate candidates

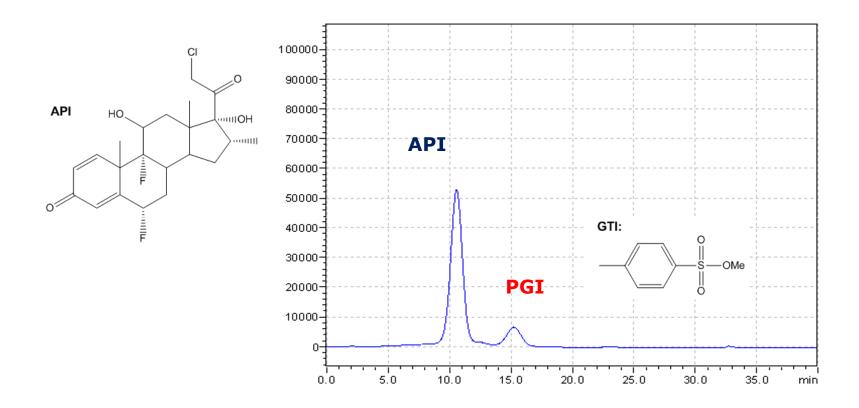


Binding curve for Diisopropylurea and API: Toluene



Arrow shows API versus PGI extraction on EAO88
= Starting point for optimized separation





Separation in LC mode on EA088
Strong retention of toxic contaminant obtained



CS 3: Conclusions

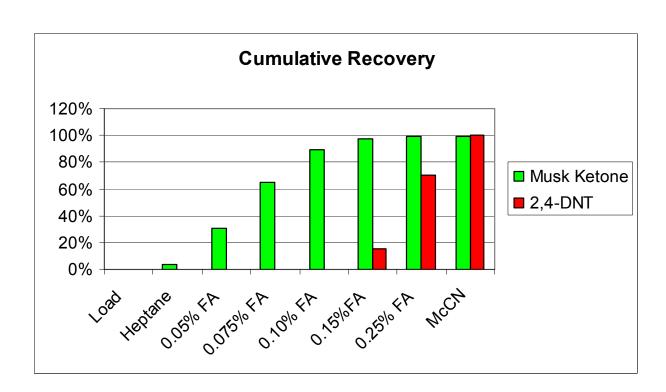
- Initial screening showed weak hits
- Additional synthesis and screen showed stronger hits
- Packing of strongest hit in HPLC mode showed satisfactory separation



Case Study 4 European consortium study



Case Study 4



Recognition of molecules carrying the same functional group (-NO₂)



Summary

- The cross-reactivity of MIPs allows selective binding of other molecules
- This 'Similarity' concept has been used to separate API's from PGI's
- All polymers identified in ExploraSep screening can be produced at pilot and/or Process scale
- ExploraSep is a powerful tool in separation method discovery



Additional Information

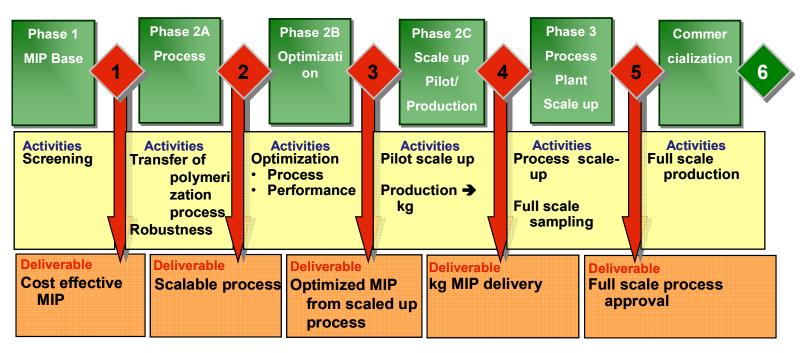


➤ How can MIPs be scaled up?



A Stage Gate approach to Production

The process path



- Applying stage-gate process for efficiency and risk reduction
- Utilising DoE (Design of Experiments) tool to attain robust product and control



Tools in materials development

- A select group of experts
- Innovative MIP chemistry → large IP portfolio
- Polymer chemistry at the research frontier
 - Novel concepts
 - New methods
- Modeling

Chemometric predictions and Monte Carlo models



Scale up – DOE Starting point

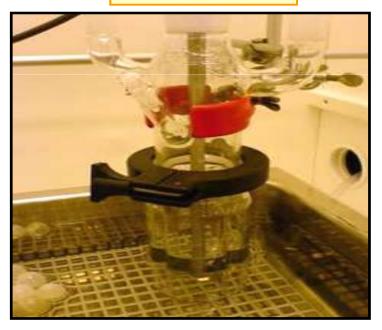
Focus

- Polymer properties
- PSD (particle size distribution)

Parallel 50ml

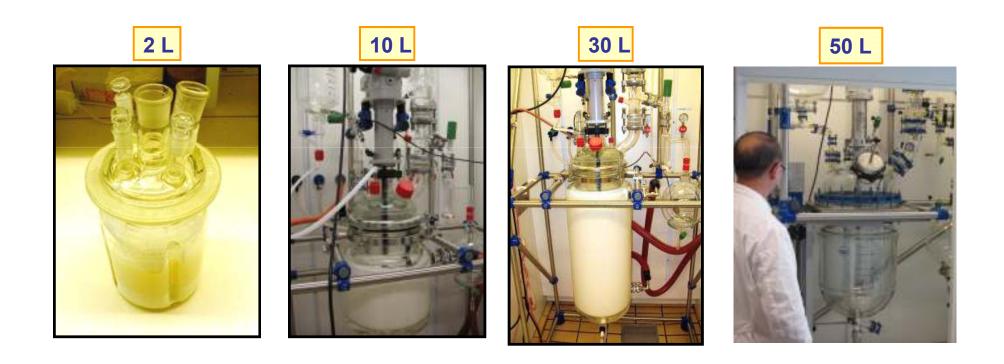


0,5 L reactors





Transfer to pilot scale ...





Example – Selective removal of nitrosamines in presence of nicotine

- To selectively remove a group of 4 nitrosamines at a total concentration of 70 ng/ml from a tobacco extract
- The structurally related nicotine should be retained in the mixture at a concentration of 700 µg/ml
- Nitrosamines are structurally related and are at 10000x lower concentration

Nitrosamines to be removed

Nicotine to be retained



Optimization of the MIP recipe

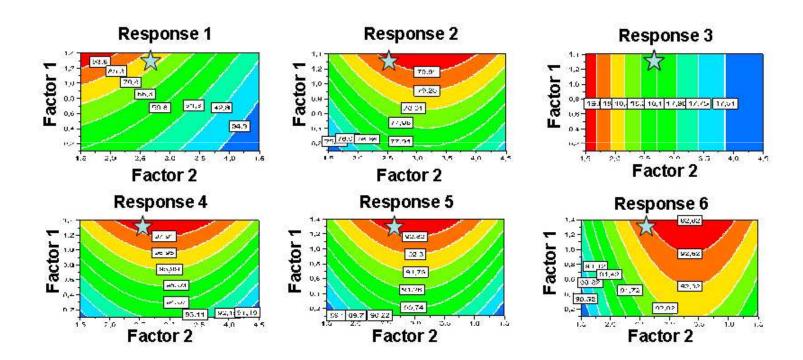
DOE computer aided optimization of the recipe

- Final DOE on "best" existing recipe
- 5 Parameters checked
- Coefficients analyzed with software MODDE
 - contour plots for the responses



The DOE Responses

DoE driven process scale-up: Optimisation/model



The Final Design: Pilot Scale (25kg)

