



Criblage de Supports Chromatographiques à Haut Débit

Innovations-Protéines-Prod

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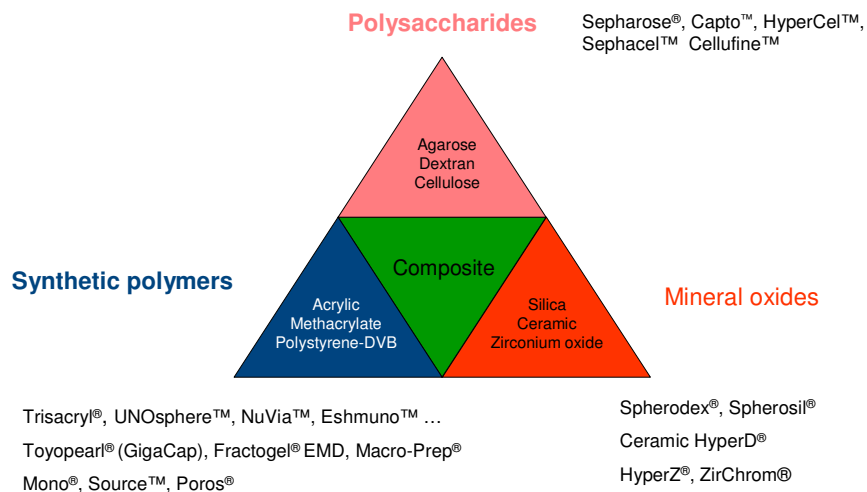
Matrices de base des supports (résines)



Les supports de chromatographie sont constitués de billes à base d'agarose, d'agarose réticulé, de polymère, de silice, de celluloses, de composites etc... (taille autour de 50-100 μm pour les applications industrielles, $<20\mu\text{m}$ pour l'analytique)



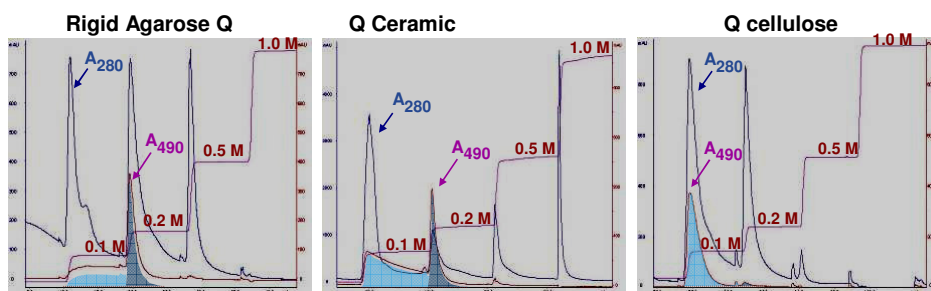
Supports de base pour la chromatographie



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Impact de la nature chimique du support de base sur la Sélectivité



Purification of rGFP from E.coli lysate on anion exchangers

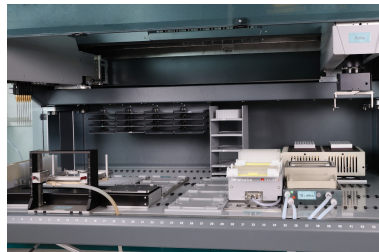
50 mM Tris-HCl pH 8.5; step elution at 0.1, 0.2, 0.5 and 1M NaCl; 0.5 cm I.D. x 5 cm at 300 cm/h (1 min RT)

Des différences de comportement chromatographique sont apportées par la nature de la matrice elle-même

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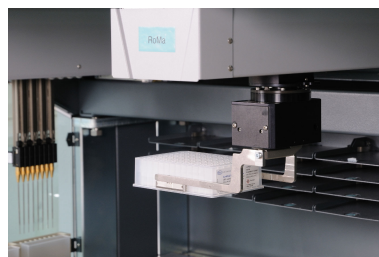


Introduction de la Robotique



Janus® BioTx Pro Workstations (PerkinElmer)

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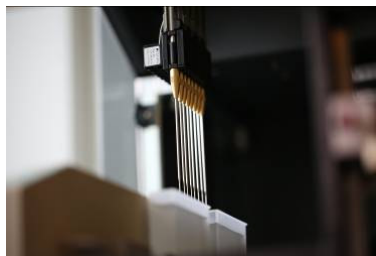
- Service d'aide au développement rapide de procédés de purification
- Différentes supports testés en parallèle: échangeurs d'ions, mixed-mode, affinité, pseudo-affinité, hydroxyapatite, HIC ...
 - Résultats en 2-3 jours grâce à des méthodes analytiques puissantes



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2013: Pall ScreenExpertSM Services Laboratory Cergy (Paris), France



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Les Méthodes d'Analyse à Haut-Débit



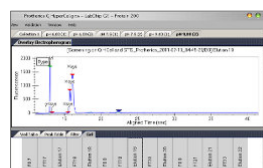
SELDI-Mass Spectrometry



Automated microfluidic
electrophoresis (LabChip®)



BLI (Bio Layer Interferometry)



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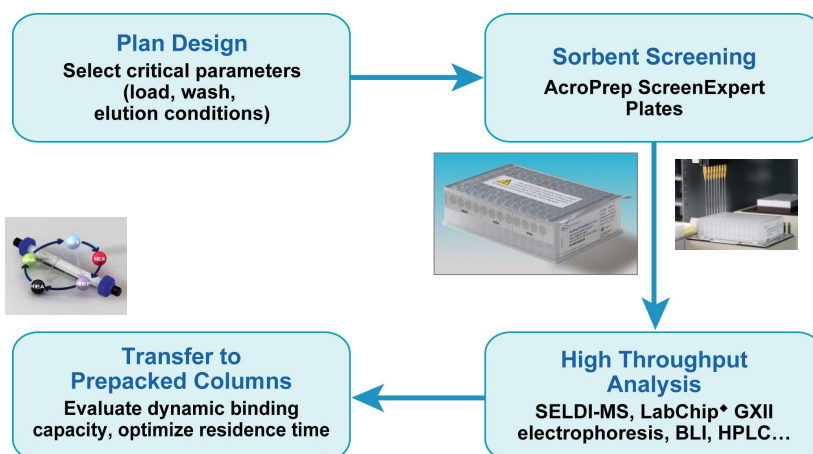


Mise au point d'un procédé de purification d'un fragment Fab d'anticorps

- Objectif: mettre au point un process de purification de Fab exprimé dans *Pichia pastoris*
- Transposable à l'échelle industrielle
- Procédé simple, avec supports robustes
- Eviter les ligands d'affinité



Screening Strategy





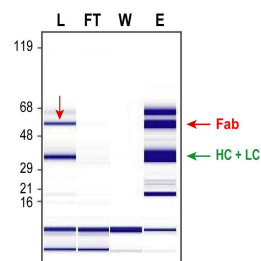
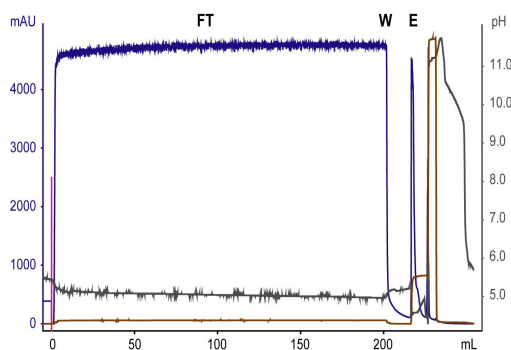
Criblage de l'étape de Capture du Fab

- **176 conditions sur 9 supports** différents testés en mode « bind/elute » comme candidats potentiels pour la capture du Fab
- Echange d'anions et and cations, Hydroxyapatite, Mixed-Mode, autres
- **Resultat : CIEX (CM) et Mixed-mode fonctionnent**
 - MEP HyperCel charge à pH 6.0 ; Fab elution à pH 5.0
 - Ou CM Ceramic HyperD, pH 5.0, elution Fab 0.15 M NaCl
- Le support CM sera retenu en raison d'une plus forte capacité dynamique (> 60 mg/mL)

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Capture step by Cation exchange: CM Ceramic HyperD® 1 mL PRC Column



Capture step performance

- Fab recovery = 96%
- Concentration = 20 times
- Purity = 40%

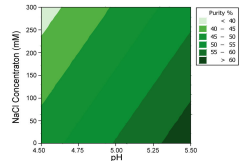
Equilibration: 50 mM Na acetate pH 5.0
Loading: 200 mL feedstock 1/2 at pH 5.0
Washing: 50 mM Na acetate pH 5.0
Elution: 50 mM Na acetate pH 5.0 + 0.3M NaCl
Flow rate: 150 cm/h – Residence Time = 3 min

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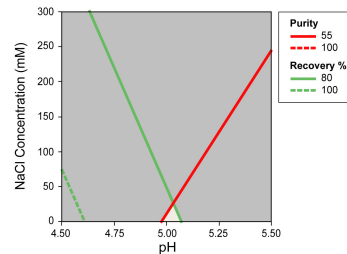
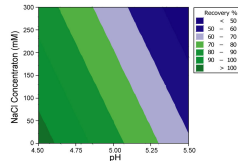


Intermediate step: Screening on AcroPrep 96 plates (Mixed Mode on MEP HyperCel)

Fab purity



Fab recovery

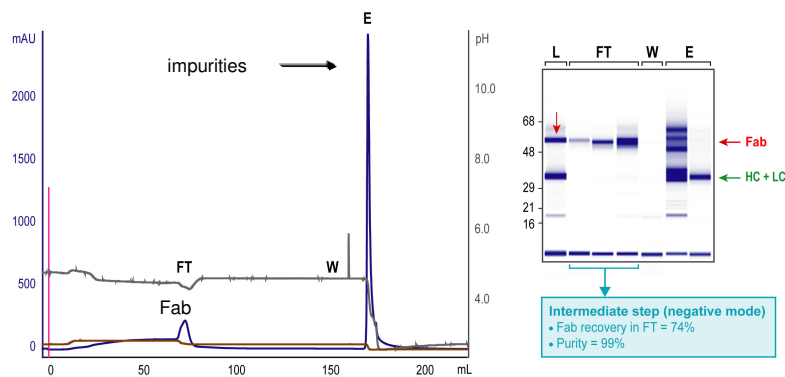


Screening of 3 pH (pH 4.5, 5.0, 5.5) and
3 NaCl conditions (0 mM, 150 mM, 300 mM)

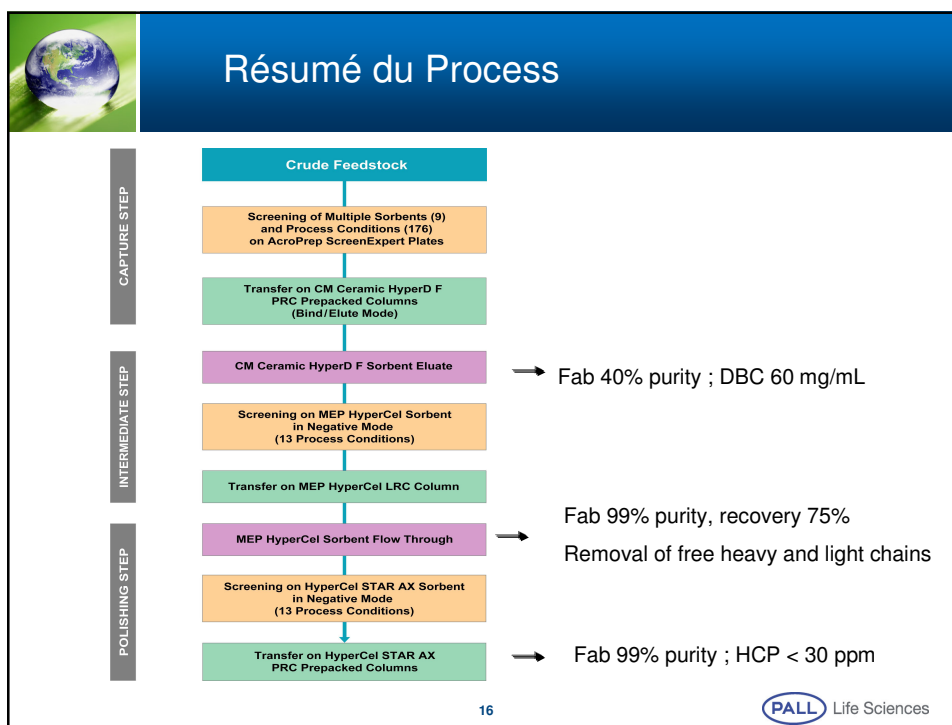
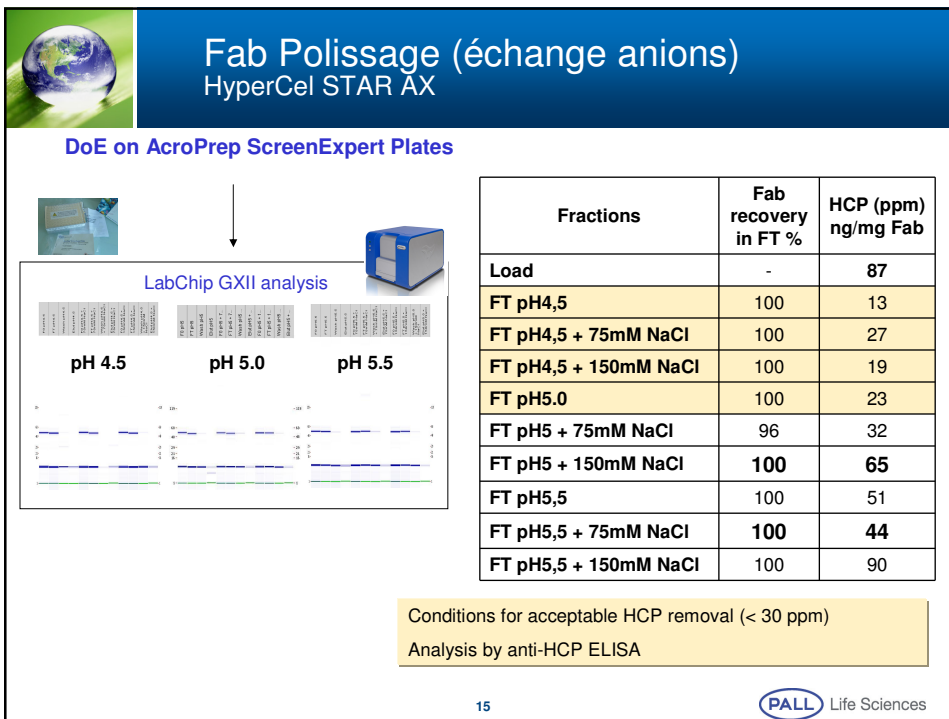
Optimal conditions on MEP HyperCel (FT mode):
Binding at pH 5.0 to 5.2, without addition of NaCl



Etape intermediaire : Mixed Mode Transfert sur colonne MEP HyperCel™ 10 mL



Equilibration: 50 mM Na acetate pH 5.2
Loading: 100 mL feedstock 1/10 at pH 5.2
Washing: 50 mM Na acetate pH 5.2
Elution: 50 mM Na acetate pH 3.0
Flow rate: 150 cm/h – Residence Time = 5 min





Conclusions



Le fragment Fab peut être purifié en trois étapes simples et transposables à la production (pas de ligand d'affinité)



Le criblage en plaques 96 puits accélère considérablement la sélection de supports (176 conditions en 2 jours)



Les paramètres optimisés doivent toujours être confirmés sur colonne (capacité dynamique)



Publications

- [1] High throughput screening of mixed-mode sorbents and optimization using pre-packed lab-scale columns for the purification of the recombinant allergen rBet v 1a. V Brenac Brochier *et al.*, **J. Chromatogr. B** (2009).
- [2] Rapid screening of purification strategies for the capture of a human recombinant F(ab')₂ expressed in baculovirus-infected cells using a micro-plate approach and SELDI-MS. J Pezzini *et al.*, **J. Chromatogr. B** (2009).
- [3] Expression optimization and purification process development of an engineered soluble recombinant mouse linker of activation of T cells using SELDI-MS. V Brenac *et al.*, **Prot. Express. and Purif.** (2006).





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