# Product characterization of pertussis Whole Cell Vaccine by mass spectrometry

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- Introduction
- Gene and protein regulation during cultivation
- Mass spectrometry
- Antigen composition as measurement of product quality



# Reasons for characterization of pertussis vaccines

- Resurgence of pertussis in populations well vaccinated with acellular pertussis vaccines triggers development of new pertussis vaccines
- Polio eradication triggers development of hexavalent DPT-Hib-HepB-IPV<sub>sabin</sub> vaccines necessitating investigation of compatibility of pertussis component
- Potency release tests are not sensitive and reproducible enough and therefore not suitable for this type of research



### **B.** pertussis culture

- Start of all pertussis vaccine productions
- Undefined intermediate product
- Product characterization required



### **Process and product characterization**





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### Genomics

Bvg-regulated virulence of *B. pertussis* 





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### **Process knowledge = product quality**

#### Culture conditions of *B. pertussis* affect the quality of WCV.

- Bvg regulated virulence factors are in general protective
- Expression of virulence factors affected by e.g. nutrient limitations, culture temperature ٠ (artificial factors like nicotine or MgSO4)



#### B. pertussis batch culture

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#### Score based on activation of 56 vag's



### Consistency = process & product knowledge

#### Steady state culture of *B. pertussis*

- MgSO4 → *bvg* down regulation → reduction in virulence factors
- Product characterization
  - 3 biorector runs
  - sampling  $\rightarrow$  T = 0, 2, 6, 12 and 24 hours
  - Plain inactivated WCV A, B, C, D and E







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### **Product characterization**





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### **Microarrays**

#### **Monitoring virulence associated genes**





### Monitoring key marker antigens

#### Selection of key marker antigens

- known protective antigens (generally virulence factors)
  - Ptx (excreted  $\rightarrow$  hardly present in WCV)
  - FHA
  - **Prn** (splitted off  $\rightarrow$  poorly present in WCV)
  - Fimbriae
  - Vag8 (virulence associated gene 8)

#### ELISA (quick scan)

- Coating serial dilutions of whole cell products
- Detection of specific antigens with MoAb's
- Goat anti Mouse IgG-HRP

#### Is this the right selection?

- Gene activation
- Protein production
- Immunoprofilling





### Identification of proteins by LC-MS





MS/MS spectra Peptides (protease fragments) of peptides Protein m/z Identified Matching peptides/ proteins in silico MS/MS pattern Peptides predicted Protein from proteolysis database from theoretical peptides m/z



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### **Quantitation of proteins by LC-MS**





### Quantitation of peptides and proteins



Up and down regulation of protein expression

	T=0	T=6	T=12	T=24
Pertactin	1.69	1.06	0.48	0.15
(P14283)				
Putative periplasmatic substrate binding protein (Q7VWX9)	0.11	1.03	1.82	2.47



### Gene and protein expression





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### % virulence proteins in WCV



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### **Relation antigen content and protection**





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# Vaccine protein composition

**2DE** 

LC-MS





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### Vaccine protein composition





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### **Proteomics whole cell based vaccines**

	WCV	OMVs	
Peptides identified	7600	_	
Proteins identified	1200	-	
Proteins quantified	332	268	
Proteins differently expressed	151	-	
% virulence factors	≈25%	50-75%	
Top 5 of known antigens	groEL, fhaB, vag8, tcfA, brkA	Vag8, brkA, tcfA, groEL,sphB1	



### Conclusions

Antigenic composition of whole cell based pertussis vaccines is largely determined by production process conditions

Key marker antigens

- expected: Ptx, FHA, Prn, Fimbriae
- determined: FHA, Vag8, BrkA, TcfA

Efficacy of pertussis vaccine seems to be related to the proportion of specific antigens in the product

### LC-MS is a powerful tool to characterize undefined vaccines like WCV



More detailed information can be requested by mail:

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