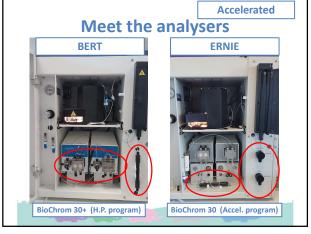


What's it all about?

- Amino Acid sample processing (in brief)
- What the Chromatography can tell us
- Pressure problems and what to do about them
- Understanding the program and how it can help us

2



Amino Acid Sample Processing

3

Processing samples (in brief)

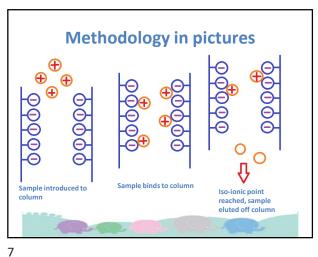
- Plasma, CSF or urine acidified and deproteinised (1:1 with IS/SSA)
- Internal standard includes Norleucine + AEC
- 44 Aminos Acids measured over 120ish minutes
- Approx 2.5-3 hours total runtime per sample
- At Alder Hey we report 24 Amino Acids
- Operator checks every chromatogram

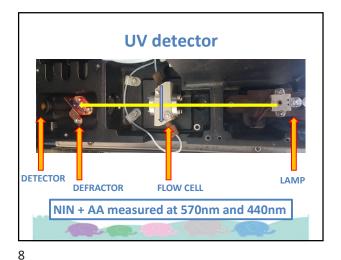
Methodology

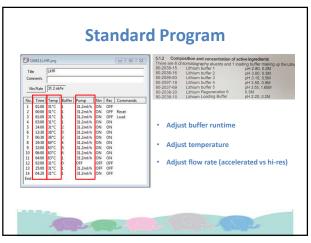
- <u>Ion Exchange Chromatography</u>
- Resin Column has -ve charged solid phase
- Acidified sample has +ve charged amino acids
- 6 mobile phase buffers of increasing pH and ionic strength
- Column temperature gradually increased
- Iso-ionic point reached AA eluted off column
- Mixed with NIN and travels to detector

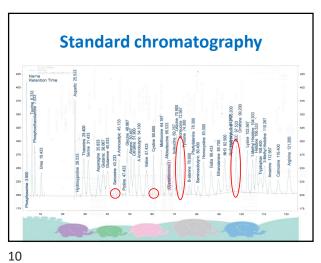
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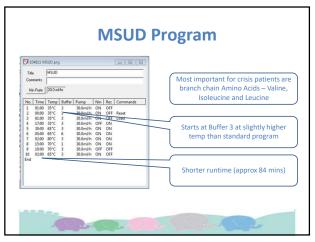
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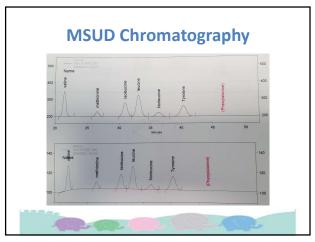


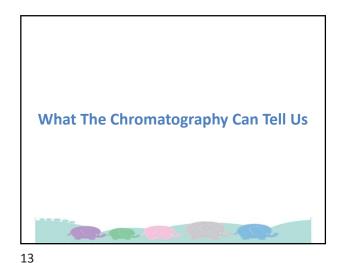








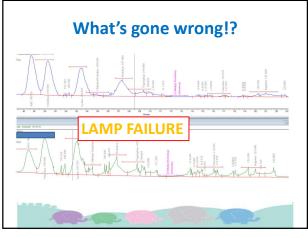


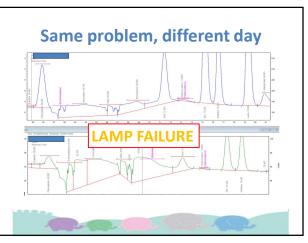


A good chromatography has...

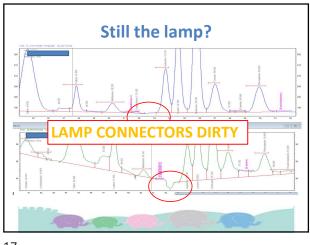
- Flat (ish) baseline
- Smooth peaks
- Symmetrical peaks
- Little no co-elution of peaks
- Clear and unobstructed AEC peak

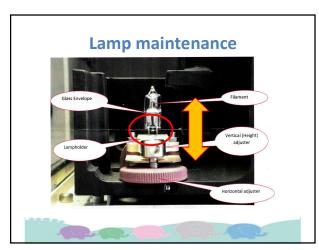
14





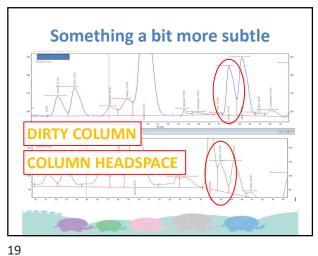
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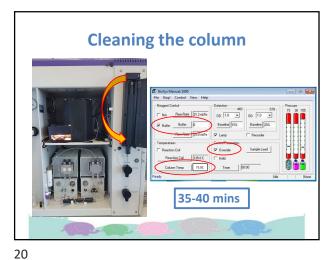


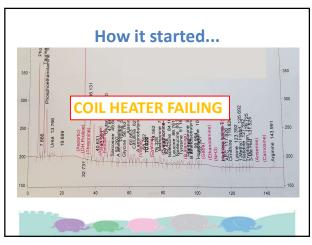


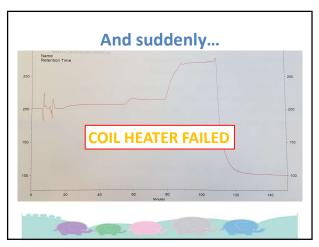
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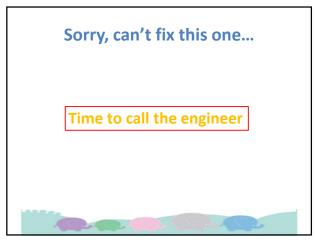
28/06/2021

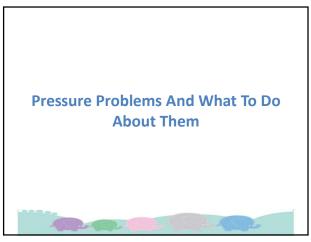


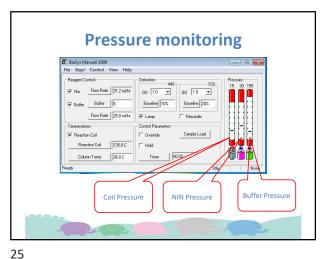


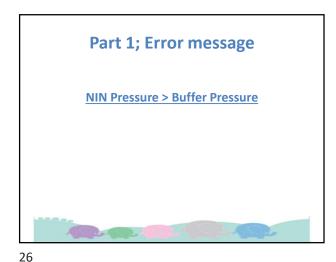


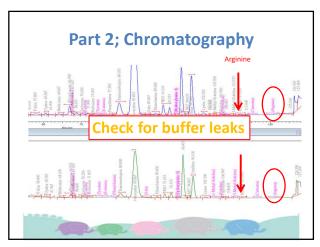




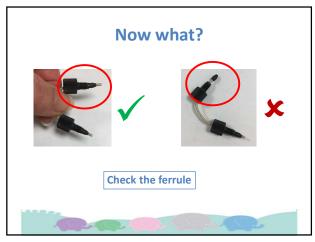


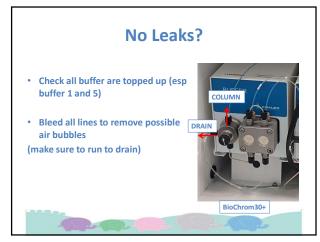


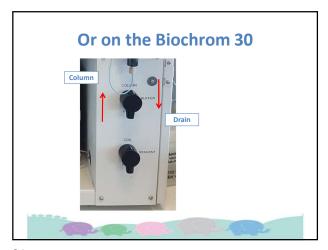


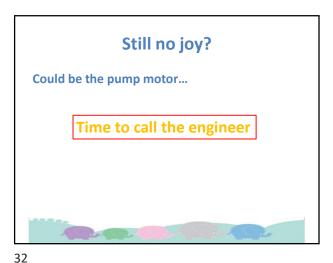




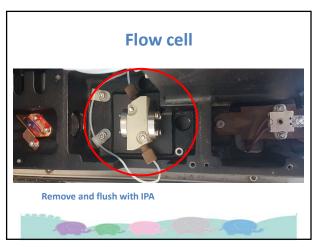




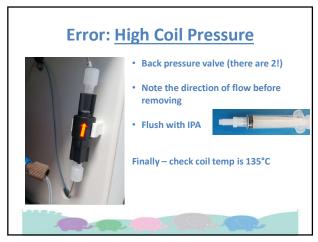


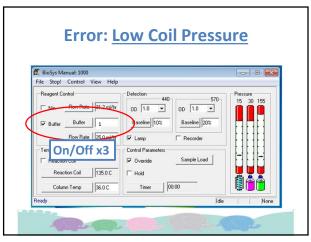




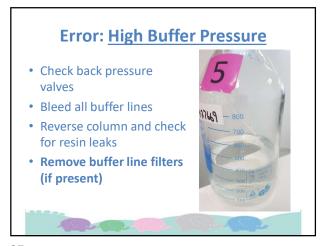


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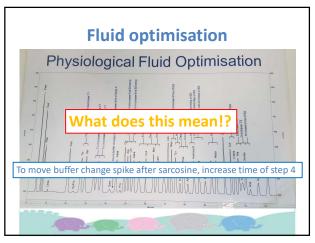


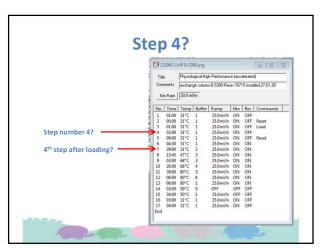


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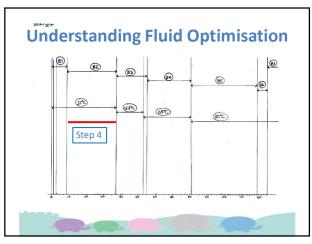


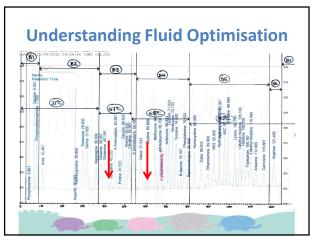




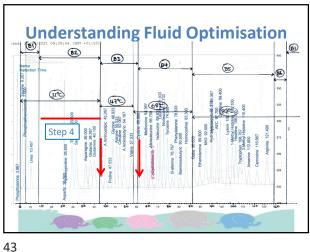


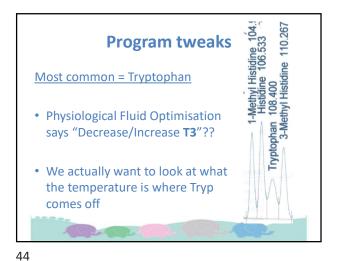
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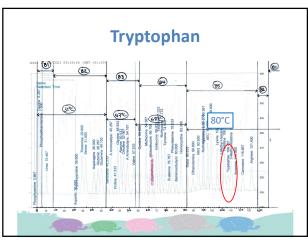


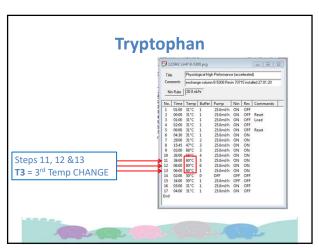


41 42









We made it to the end! • Thanks for listening, any questions?