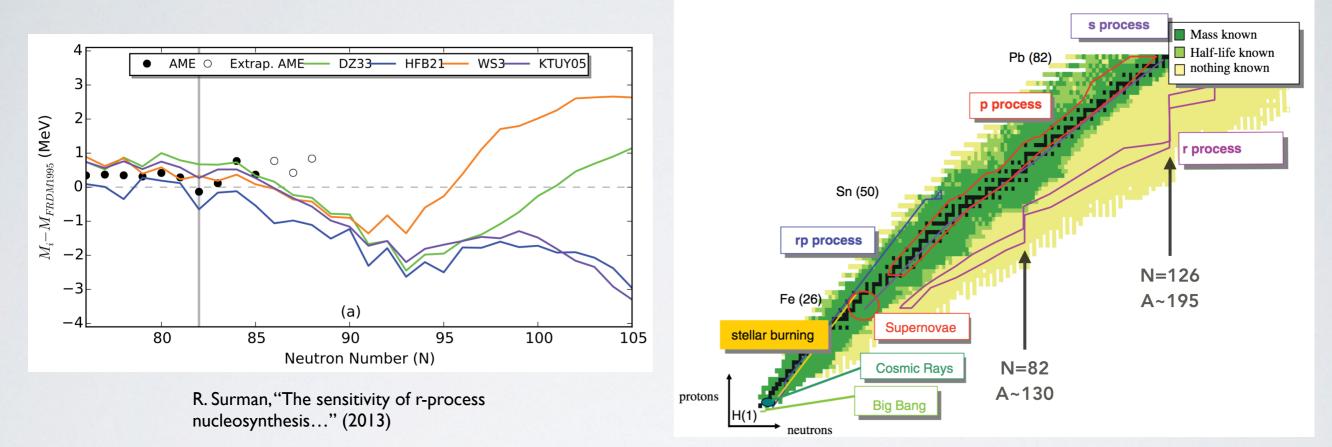
MR-TOF DEVELOPMENT AT THE UNIVERSITY OF NOTRE DAME

James Kelly

CS Workshop, GSI 21-24 March, 2016

OUTLINE

- Rapid neutron capture process
- N=126 factory at Argonne
- Notre Dame MR-TOF
- Control system



K. Blaum / Physics Reports 425 (2006) 1-78

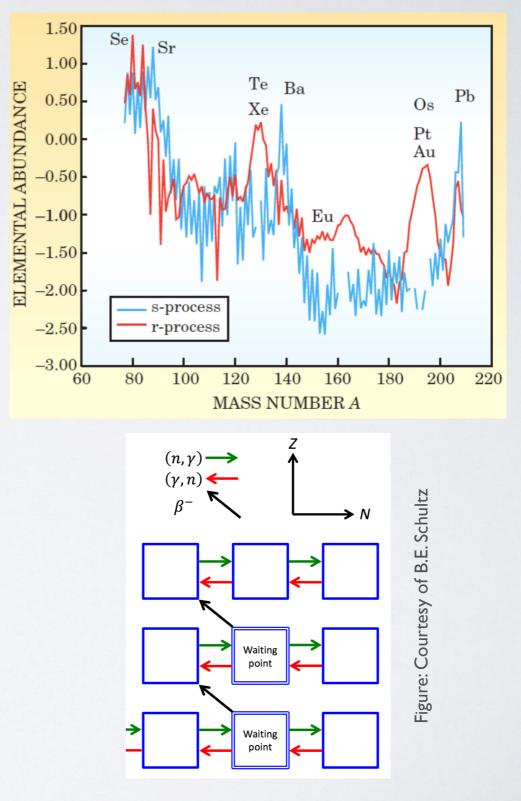
WHY MEASURE MASS?

Deformation Shell structure evolution **R-process** Unrefined models Limits of stability Halos

R-PROCESS ABUNDANCES

4

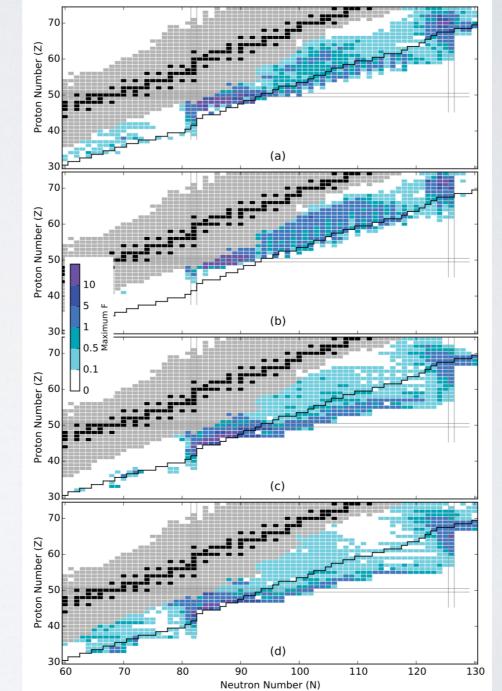
- New paths blazed in neutron-richness
 N capture outpaces photodissociation
- stellar nucleosynthesis abundance peaks seen around magic numbers
 - N=50 (A≈80): Se, Ge
 - N=82 (A≈I30):Te, Xe
 - N=126 (A≈195): Os, Ir, Pt
- Waiting point nuclei have $(n, \gamma) < ->(\gamma, n)$ equilibrium
- Nuclei at these bottlenecks are shortlived (10s of ms or less)



5

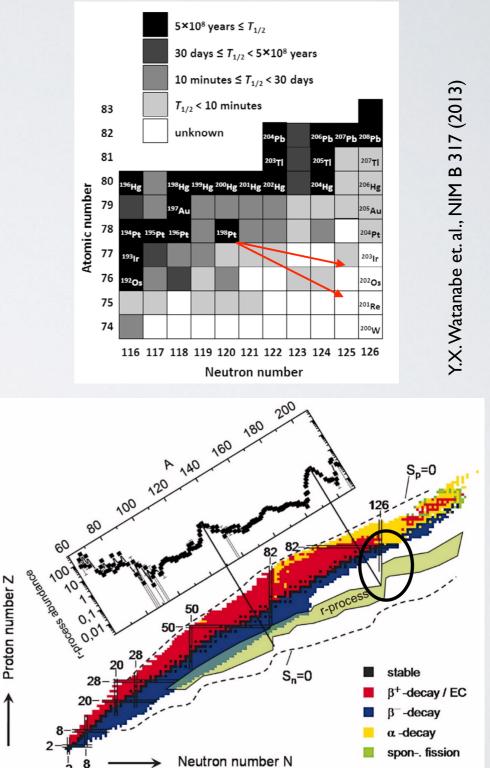
SENSITIVITY STUDIES

- Compare a baseline simulation to thousands of others, each with one piece of nuclear data varied.
 - mass, N capture cross section, ß-decay rates, ß-delayed N emission, etc.
- Mass has greatest impact on isotopic abundances.
 - especially so around closed shells due to their higher abundances.

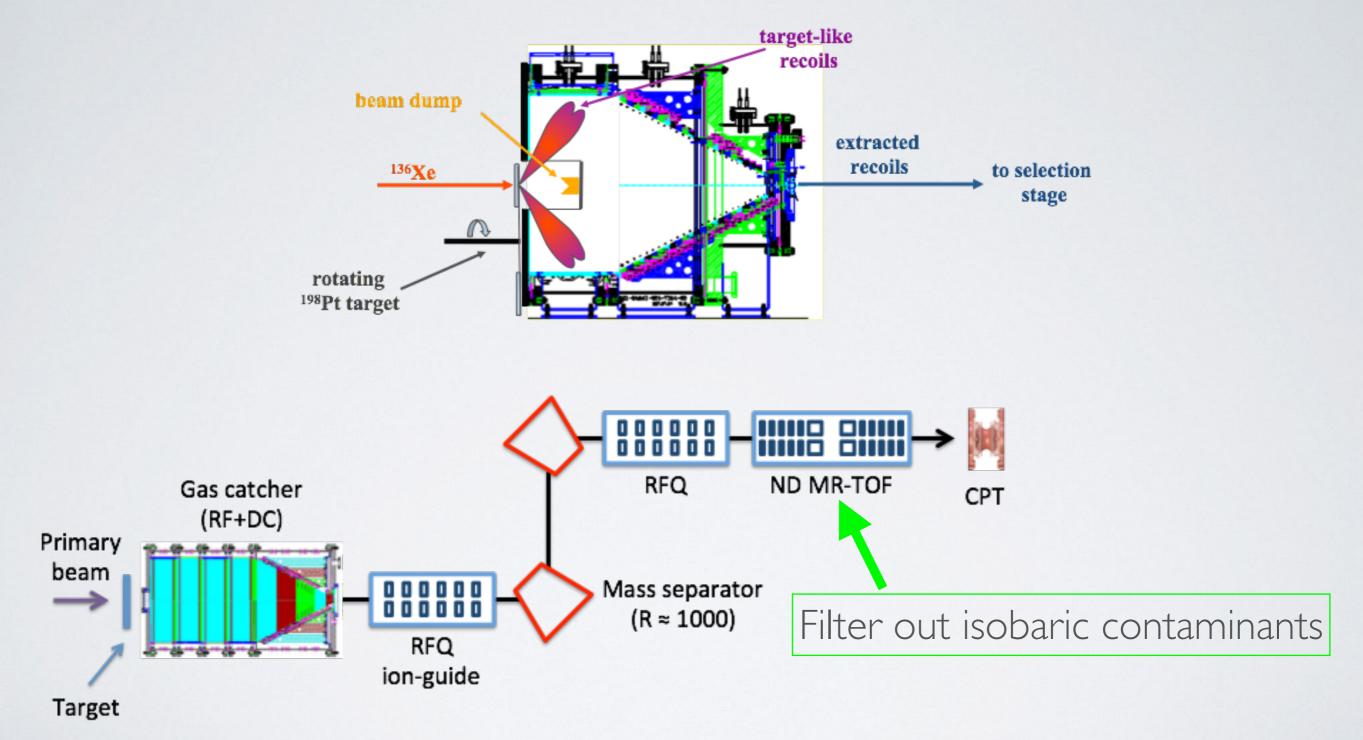


N=126 FACTORY USING ATLAS

- Argonne National Lab will house the MRTOF as part of a new facility to access N=126 nuclei.
- Produced by multi-nucleon transfer reactions (deep inelastic collisions)
 - fragmentation difficult
 - fusion, fission impossible
- I36Xe on I98Pt at <I0 MeV/A shows promise... produce nuclei about which nothing is known!



ACCESSING N=126 AT ARGONNE



MR-TOF PRINCIPLES

- Separates species (including isobars/ isomers) by mass/ charge ratio.
- Hundreds of rounds trips, measurement time ~10 ms.
- $R > 10^{5}$ possible.

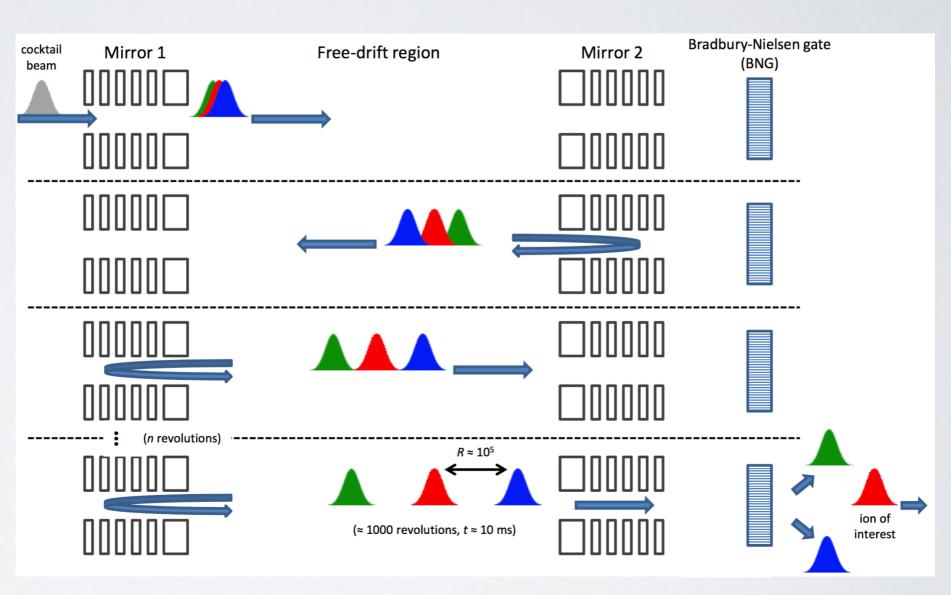
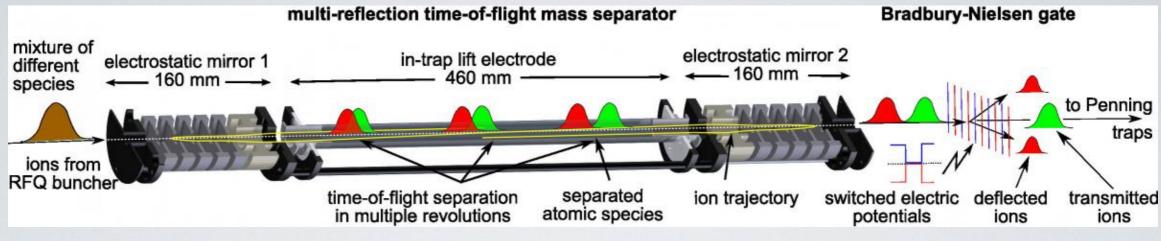


Figure: Courtesy of B.E. Schultz



CERN PH Newsletter (Mar 23, 2013)

- Trapping: In-trap lift method
 - No switching of end mirrors, which would negate the inherent precision and speed of the system.
 - Trapped energy is independent of Injection energy, and so can be tuned.

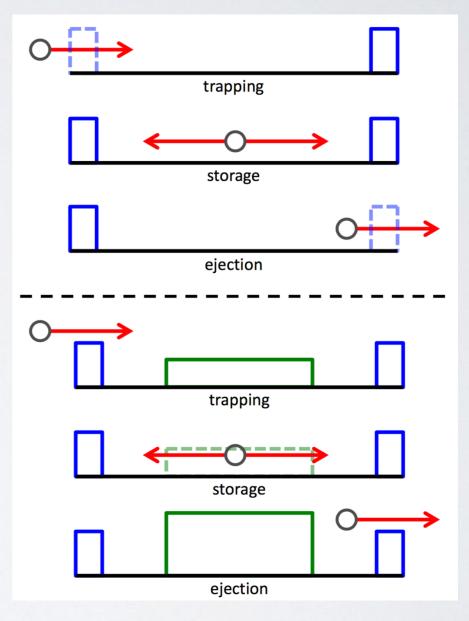
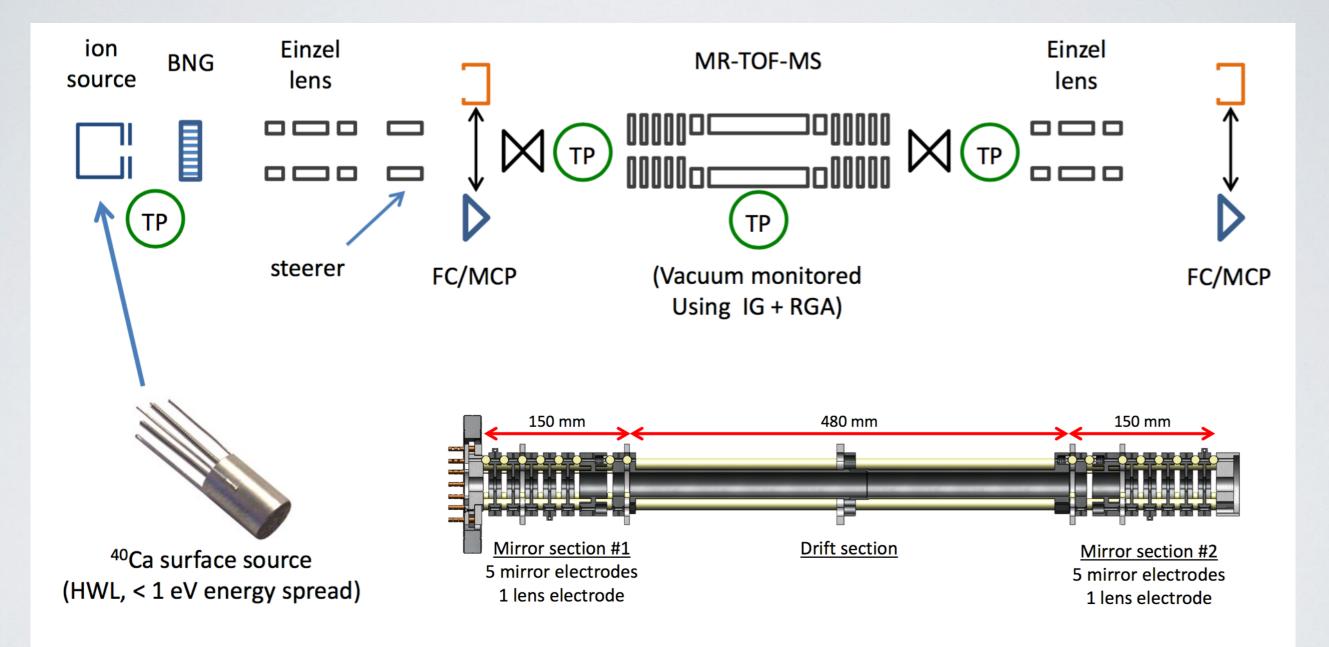


Figure: Courtesy of B.E. Schultz

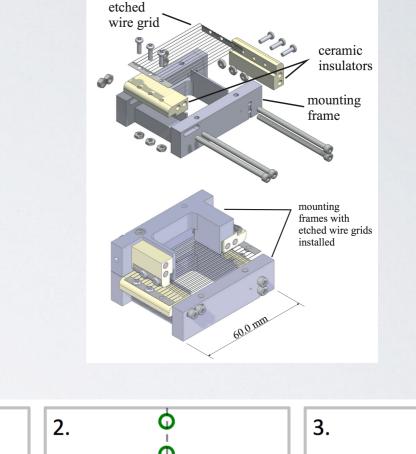
OFFLINE TEST SETUP

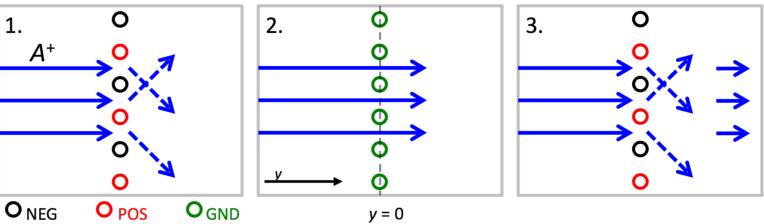


Figures: Courtesy of B.E. Schultz

SIMULATIONS AND OPTIMIZATION

- Ion bunch creation using Bradbury-Nielsen Gate
 - SIMION 8.1
 - A = 40, q = +1, E = 4600 +/- 0.3 eV (typical spread of surface ion source), 3mm beam radius
 - 100 ns bunch chopped out of "continuous" beam



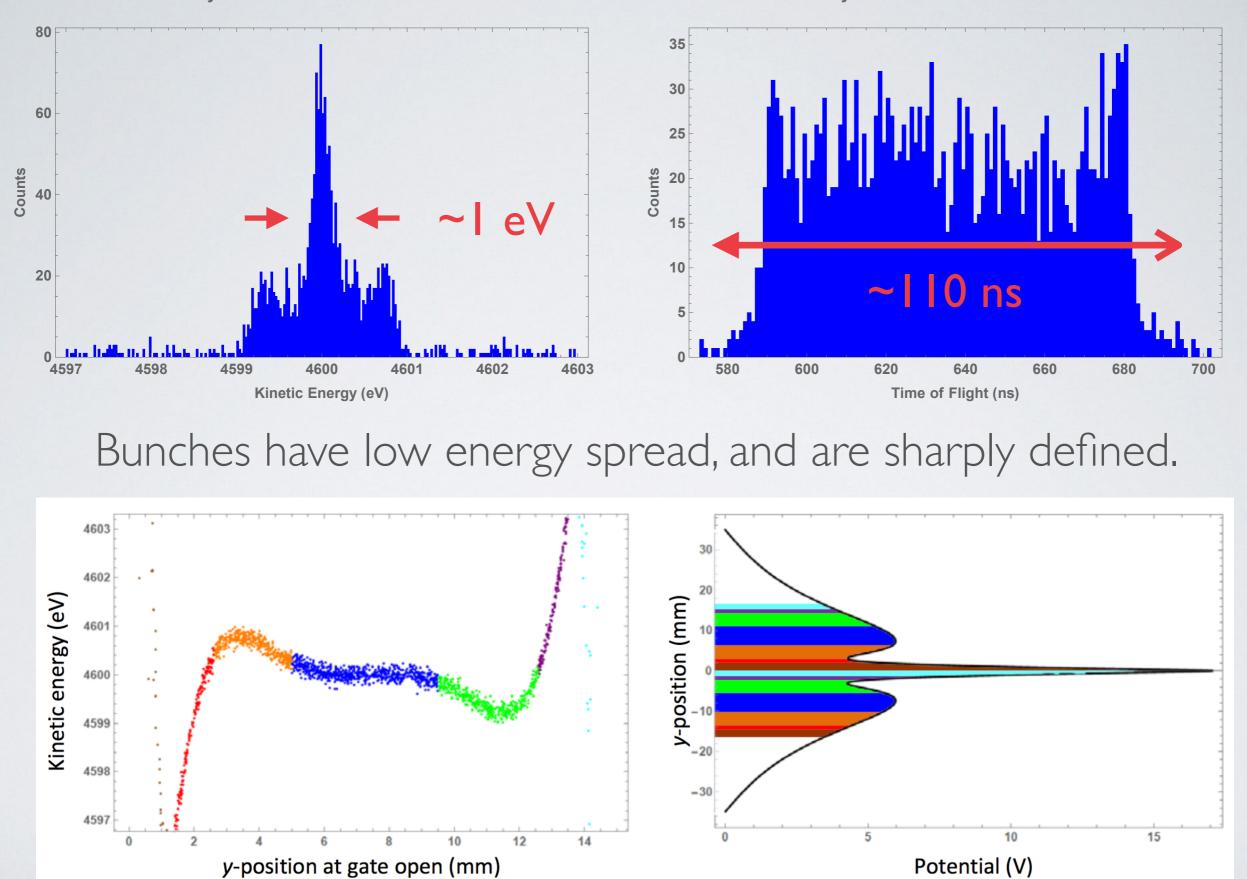


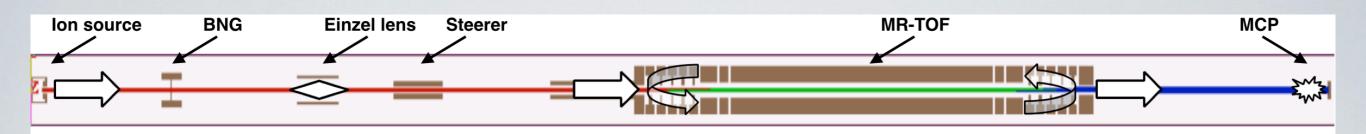
$$\tan \alpha = \frac{\pi}{2 \ln \left(\cot \left(\frac{\pi R}{2 d} \right) \right)} \frac{V_{wire}}{E_{kin}/q}.$$

Asymmetric Bias: +1100/-1060 V

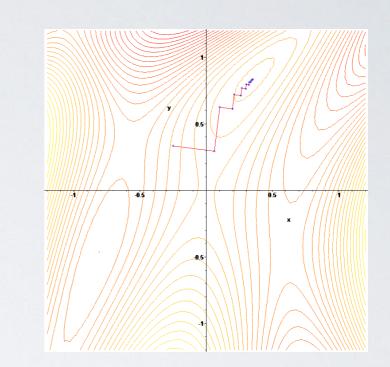
y-position at gate open (mm)

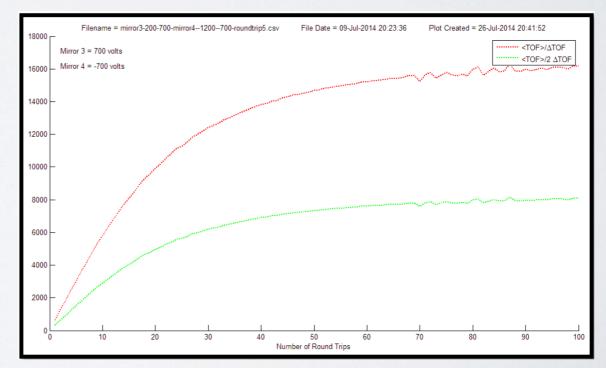
Asymmetric Bias: +1100/-1060 V

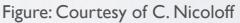


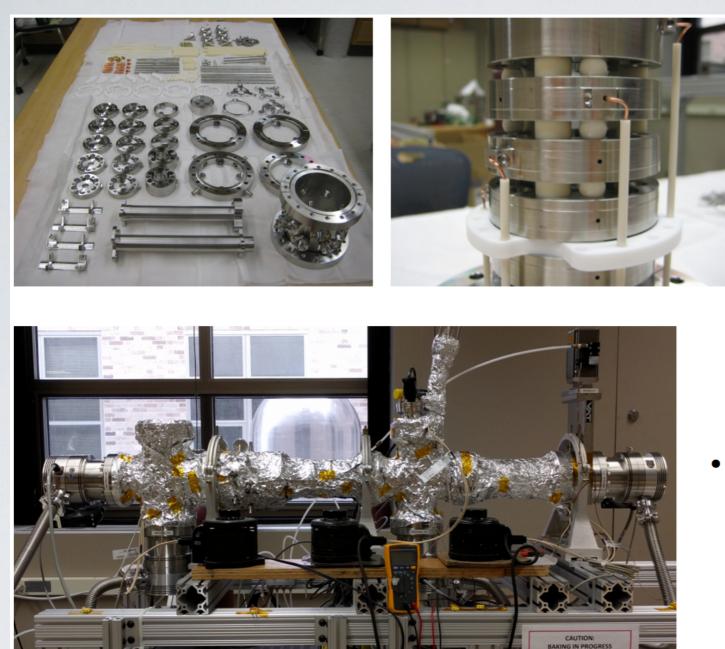


- Offline test setup simulation and MR-TOF optimization
 - adding injection and optics builds on earlier sims which started ions at MR-TOF center.
 - "gradient ascent" procedure written in lua automates the optimization process.











All components designed, cleaned and assembled for UHV compatibility



CONSTRUCTION

HOT DO NOT TOUCH

CONTROL SYSTEM & DAQ

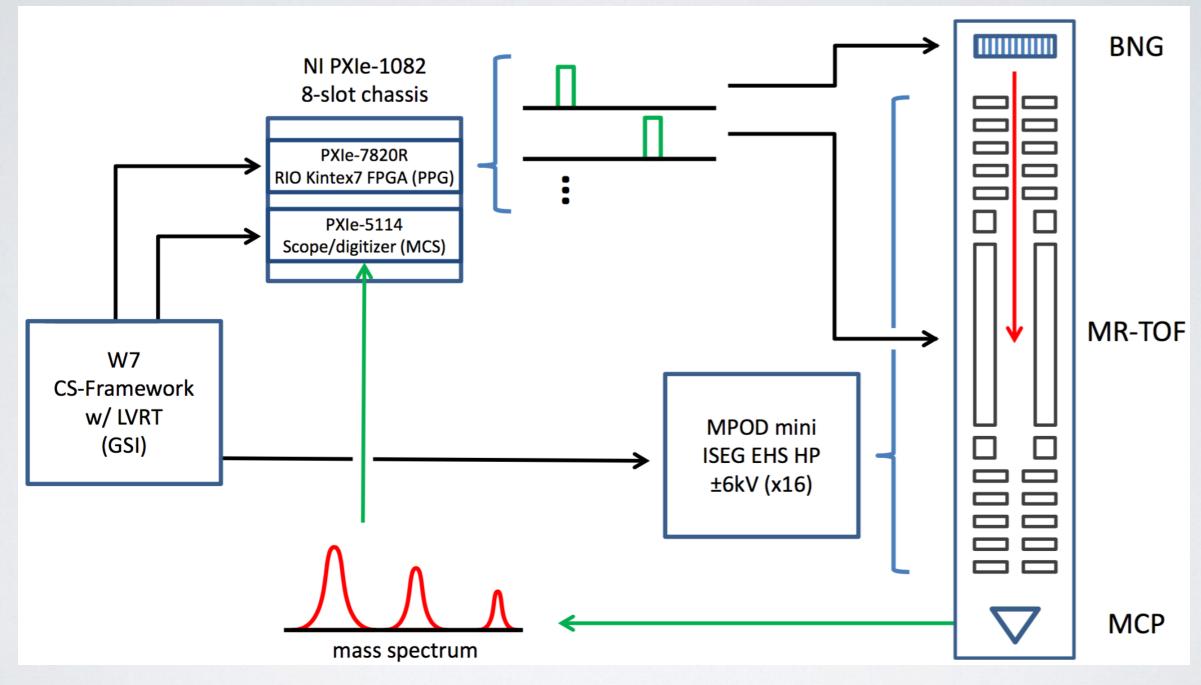


Figure: Courtesy of B.E. Schultz

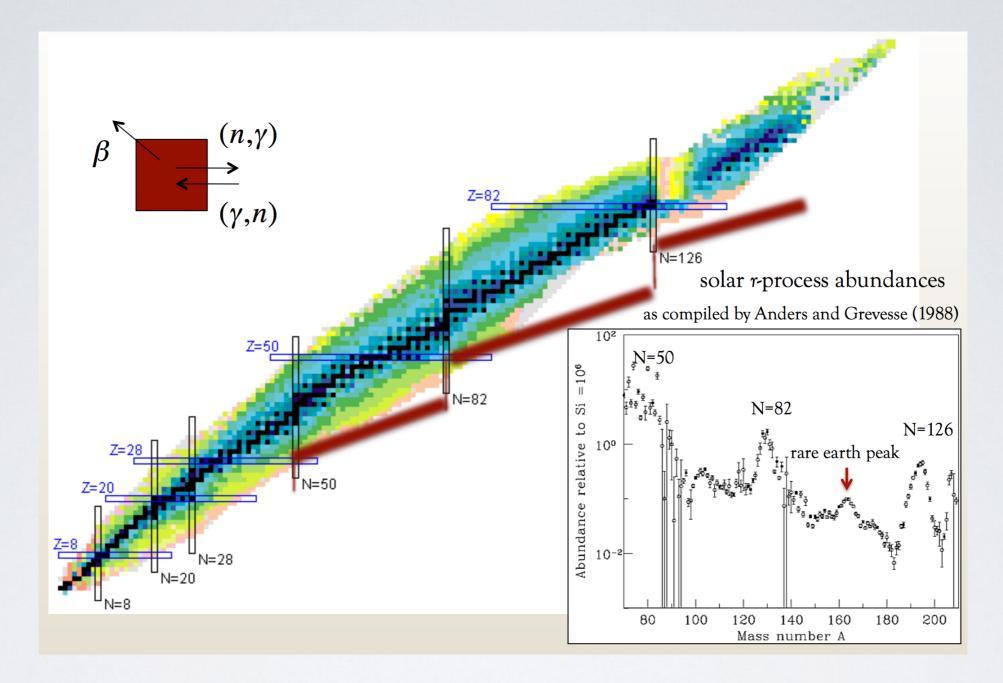
SUMMARY AND OUTLOOK

- R-process produces half of the elements heavier than iron, yet is poorly understood
- N=126 factory at Argonne National Lab will permit new measurements with the help of our MR-TOF.
- MR-TOF assembled.
- Simulations to optimize its performance are ongoing.
- HV power supplies on order. Control systems in development...
- Offline commissioning this summer. Move to ANL at beginning of 2017.

ACKNOWLEDGEMENTS

- Maxime Brodeur (Notre Dame)
- Brad E. Schultz (Notre Dame)
- Catherine Nicoloff (Wellesley College)
- Dennis Neidherr (GSI)
- Ryan Ringle (NSCL)
- Martin Eibach (NSCL)

Extra Slides



R. Surman, "The sensitivity of r-process nucleosynthesis..." (2013)

Production rate for N=126 nuclei

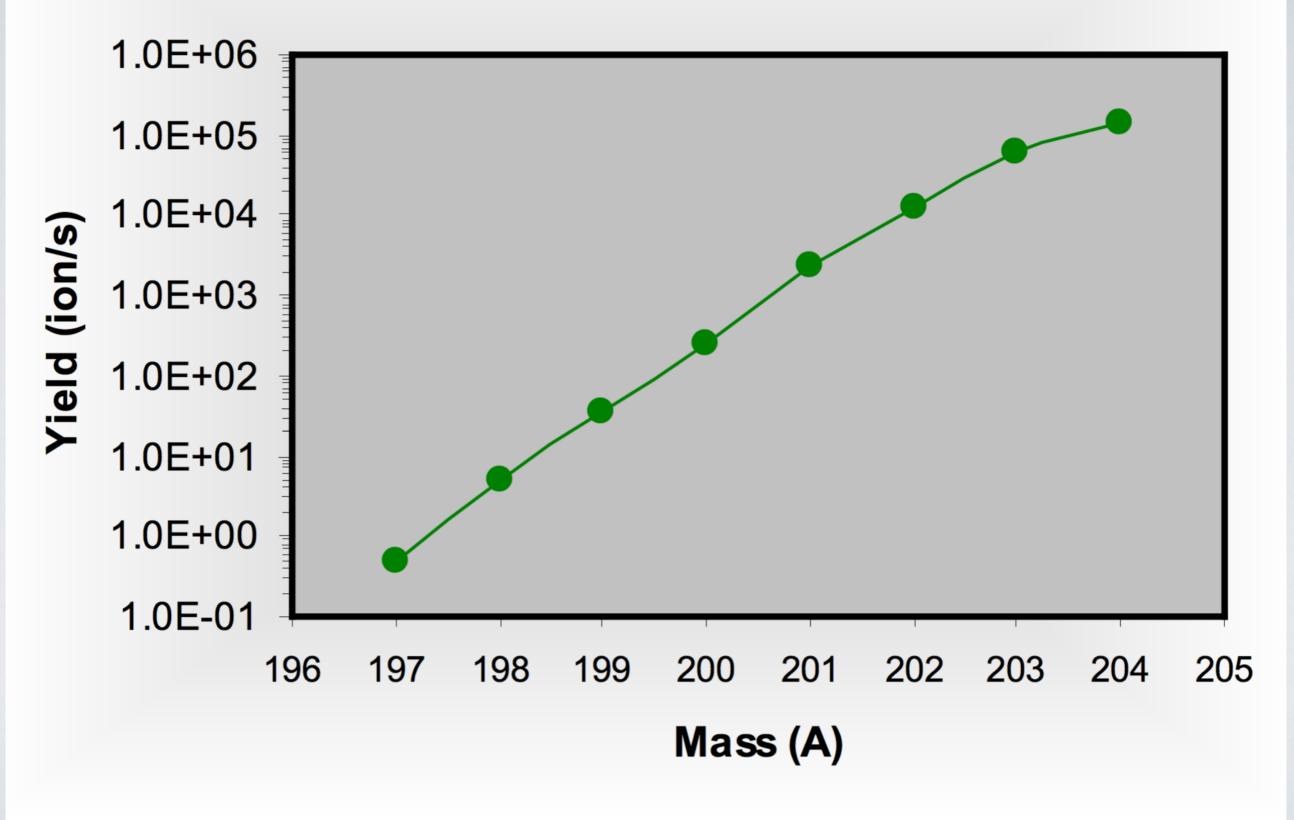
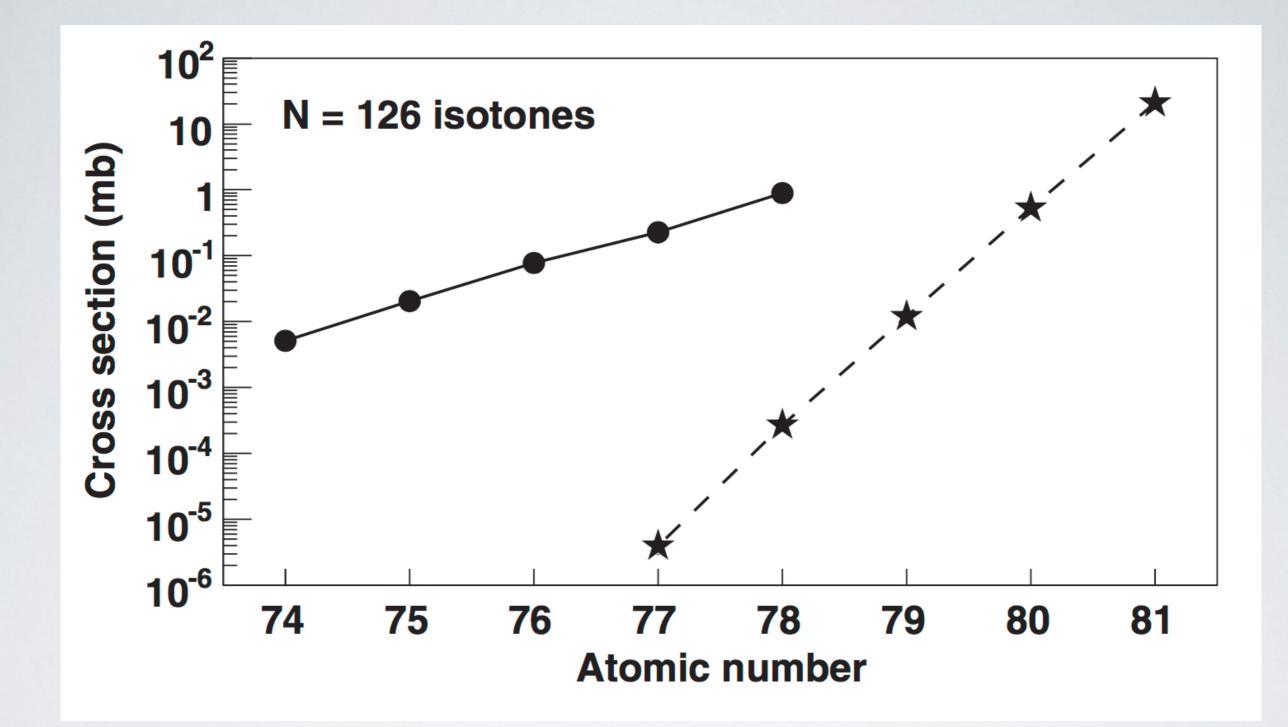


Figure courtesy of G. Savard.



MERITS OF MULTI-REFLECTION TIME-OF-FLIGHT MASS SPECTROGRAPH

- Used for measurement, diagnostics, beam purification, bunching.
 - Short measurement times (~ms) and large mass range (better than Penning trap mass spec.).
 - Requires very few ions (hundreds).
- Produces pure samples out of contaminated beam.
 - More accepting of large, contaminated bunches than Penning traps. Measure multiple species in single spectrum.
 - Accepts larger energy spread in incoming bunches due to high cycling rate.
- Extends time-of-flight (TOF) indefinitely while remaining small in size.
 - high sensitivity and resolving powers (R=m/ Δ m \geq 10⁻⁵). Comparable uncertainties to PTMS.
 - better than dipole separators (R < 5000) and HRS (R \approx 20,000)
- Resolves isobars & isomers.