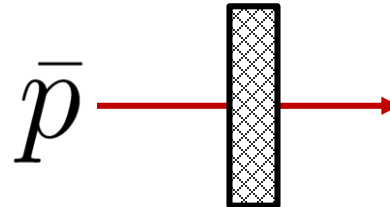
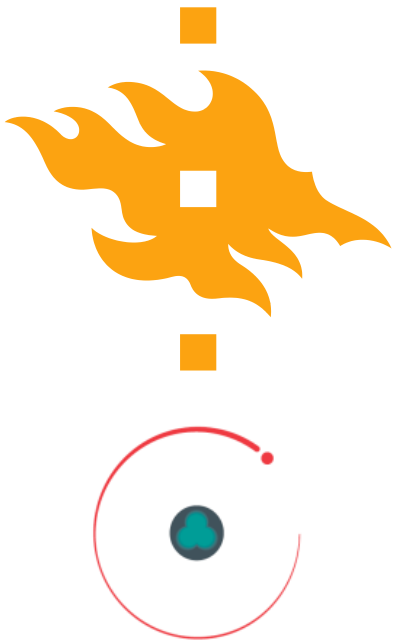


Molecular dynamics simulations of antiproton transmission through energy degrading foils



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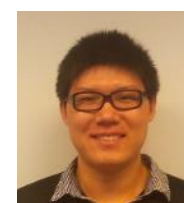
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Nuclear materials



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M.Sc. Alvaro Lopez
Surface ripples



M.Sc. Ekaterina Baibuz
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M.Sc. Mihkel Veske
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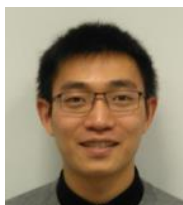
M.Sc. Simon Vigonski
Particle physics mat'ls



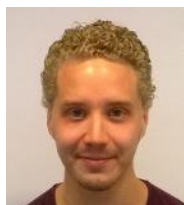
M.Sc. Henrique Muinoz
Swift heavy ions



M.Sc. Jesper Byggmästar
Fusion reactor mat'ls



M.Sc. Jian Liu
Carbon nanomaterials



B.Sc. Emil Levo
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M.Sc. Jyri Lahtinen
Machine learning

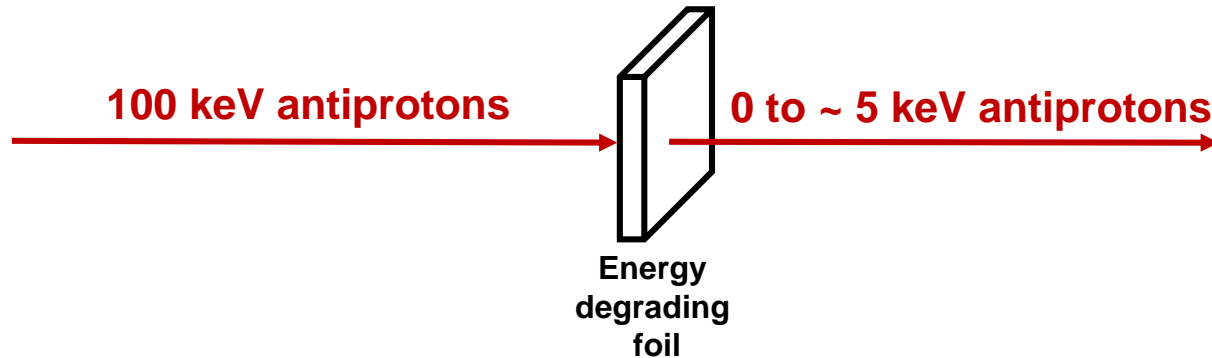


M.Sc. Anton Saressalo
Arcing experiments

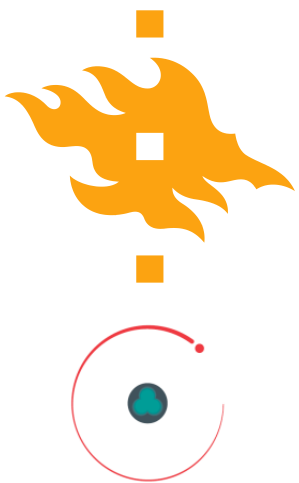


B.Sc. Jonna Romppainen
Particle Physics Mat'ls

Motivation: how to calculate the slowing down of antiprotons from ELENA in energy degrading foils?

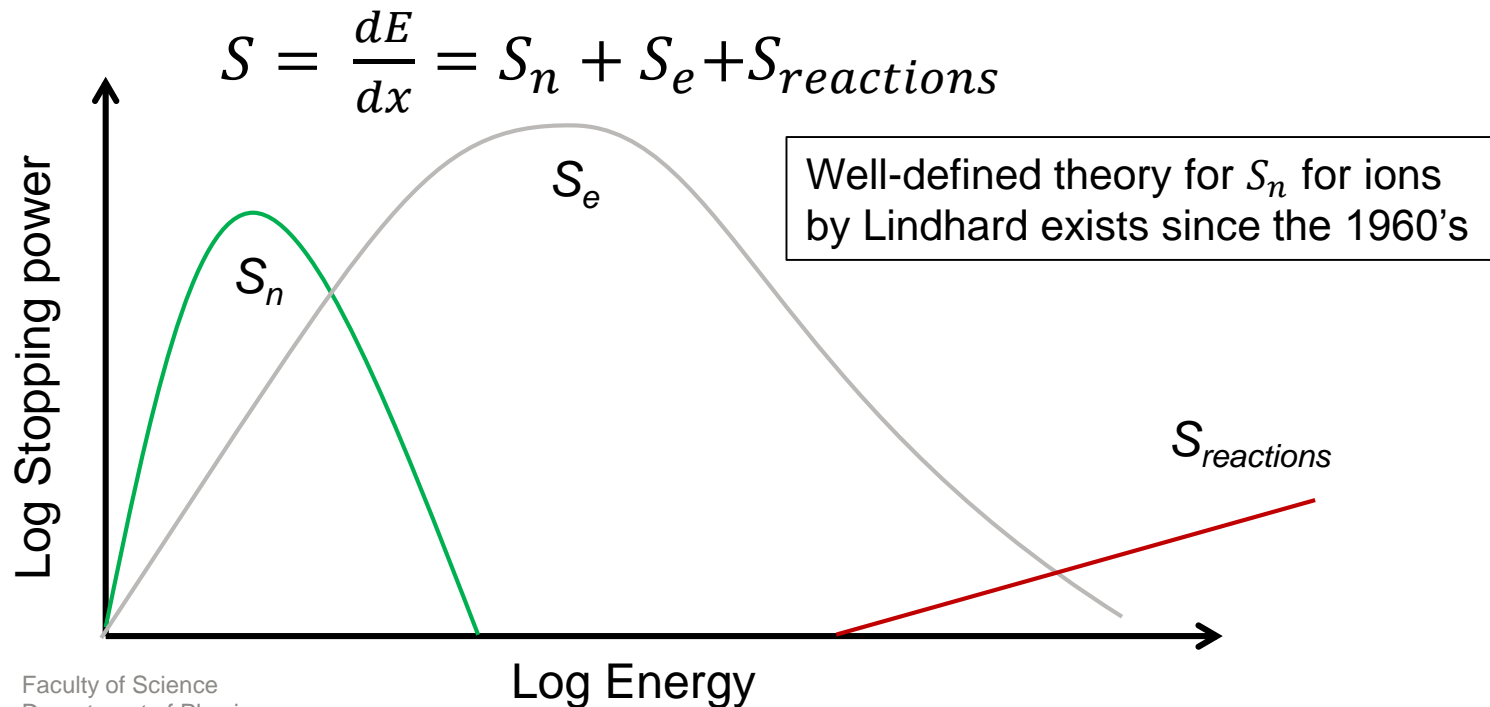


- Key question for improving antiproton yields for \bar{H} production: how to maximize fraction of 0 – 5 keV antiprotons coming through foil?
 - Foil thickness, material?
- To be able to calculate this, we need to be able to calculate **the slowing down** of antiprotons in materials



Basics of ion/anti-ion slowing down

- The slowing down of energetic ions in materials is determined by the **stopping power**
 - Sometimes also called the “linear energy transfer” LET
- Stopping power S is divided into nuclear and electronic components and nuclear reaction parts





Nuclear stopping power of antiprotons?



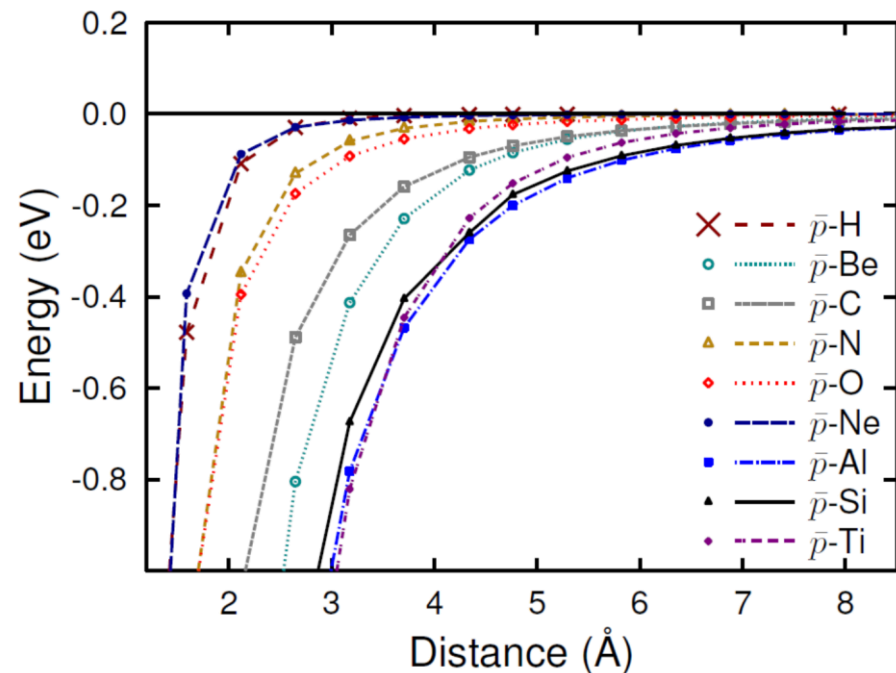
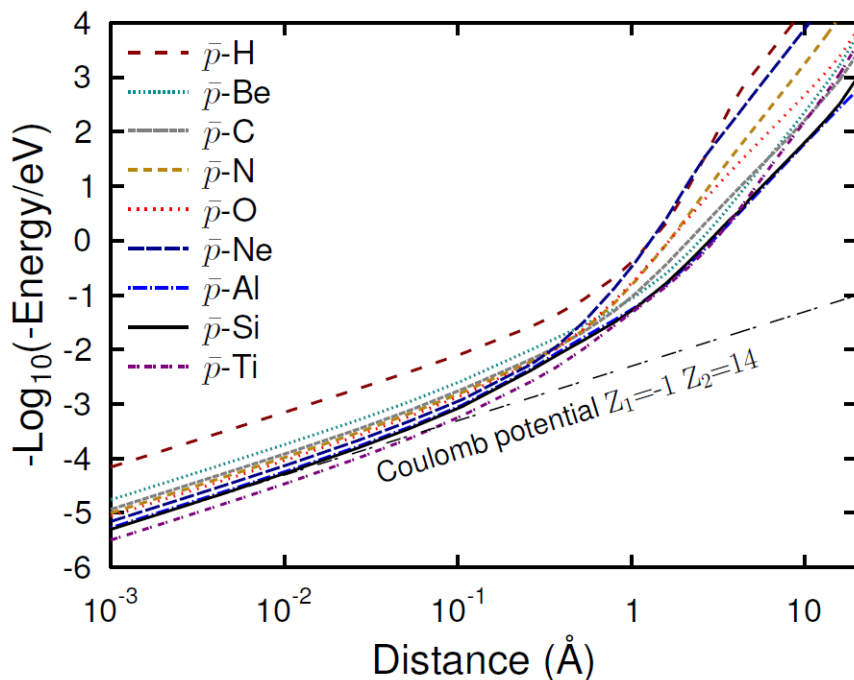
- The electronic stopping power S_e of antiprotons is known from experiments and theory in several materials
- Since we are now interested in energies < 5 keV, also the nuclear stopping power S_n may be of interest

However, turned out nobody ever determined the nuclear stopping power of antiprotons!

- Good enough motivation for doing it now
 - *Hence we basically repeated last 60 years of nuclear stopping theory for antiprotons*

Determining nuclear stopping power of antiprotons, 1.

- Step 1: determine interatomic interaction between antiprotons and atoms
- Quantum chemistry (Hartree-Fock/MP2) with Turbomole code to get total energy of antiproton – atom systems as function of distance
- DFT also tested, but gave suspicious results

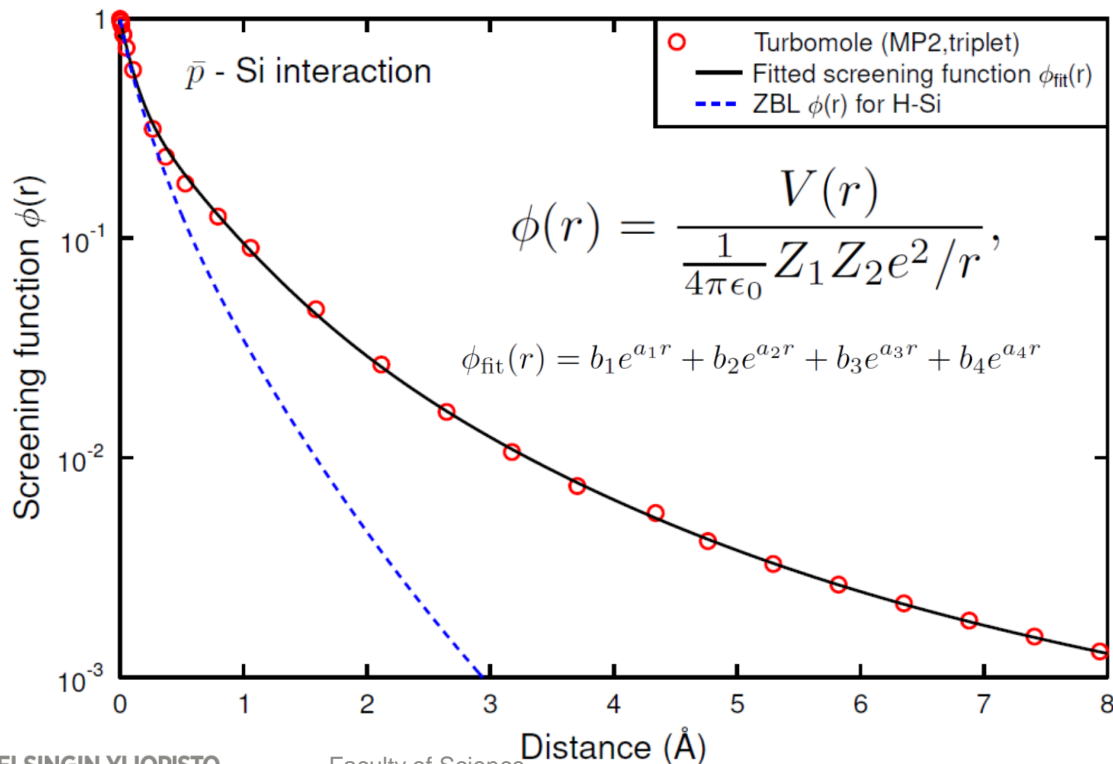




Determining nuclear stopping power of antiprotons, 2.



- Step 2: To enable practical calculations at any interatomic distance, fit screened Coulomb potential to the data
- Done for 9 elements: H, Be, C, N, O, Ne, Al, Si, Ti



| El. X | | X_1 | X_2 | X_3 | X_4 |
|-------|---|-------------|-------------|------------|-------------------------------|
| H | b | 10.688 | 0.000319357 | 0.00597013 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -2.60175 | -0.231507 | -0.689763 | -2.75173 |
| Be | b | 0.00466484 | 0.525641 | 0.0277569 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.217729 | -1.11801 | -0.48179 | -7.7278 |
| C | b | 0.0380302 | 0.318996 | 0.638666 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.612817 | -11.2902 | -1.84257 | -0.193946 |
| N | b | 0.0181678 | 0.765059 | 0.216198 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.824271 | -2.52968 | -14.7289 | -0.242982 |
| O | b | 0.0264826 | 0.790077 | 0.182093 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.847469 | -2.906 | -17.2312 | -0.200413 |
| Ne | b | 0.000113387 | 0.00617107 | 0.116765 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.253027 | -0.994693 | -24.0586 | -3.74468 |
| Al | b | 0.00128954 | 0.0254584 | 0.276114 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.097406 | -0.419499 | -1.22892 | -7.55095 |
| Si | b | 0.00547494 | 0.360324 | 0.0913726 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.19811 | -1.99426 | -0.79267 | -9.71705 |
| Ti | b | 0.00226208 | 0.574763 | 0.135421 | $(1 - b_1 - b_2 - b_3 - b_4)$ |
| | a | -0.247931 | -3.83816 | -0.924926 | -18.5785 |



Determining nuclear stopping power of antiprotons, 3.



- Step 3: Use classical scattering theory of binary collisions to calculate energy loss in binary collisions and by integrating this, get nuclear stopping power

$$S_n(E_0) = N \int_0^\infty T(E_0, b) d\sigma = 2\pi \int_0^\infty T(E_0, b) b db$$

(b is the impact parameter of the binary collision)

- Note that since the potentials are attractive, the trajectories are not necessarily hyperbolic: binary collision approximation cannot be used, needed to use molecular dynamics!

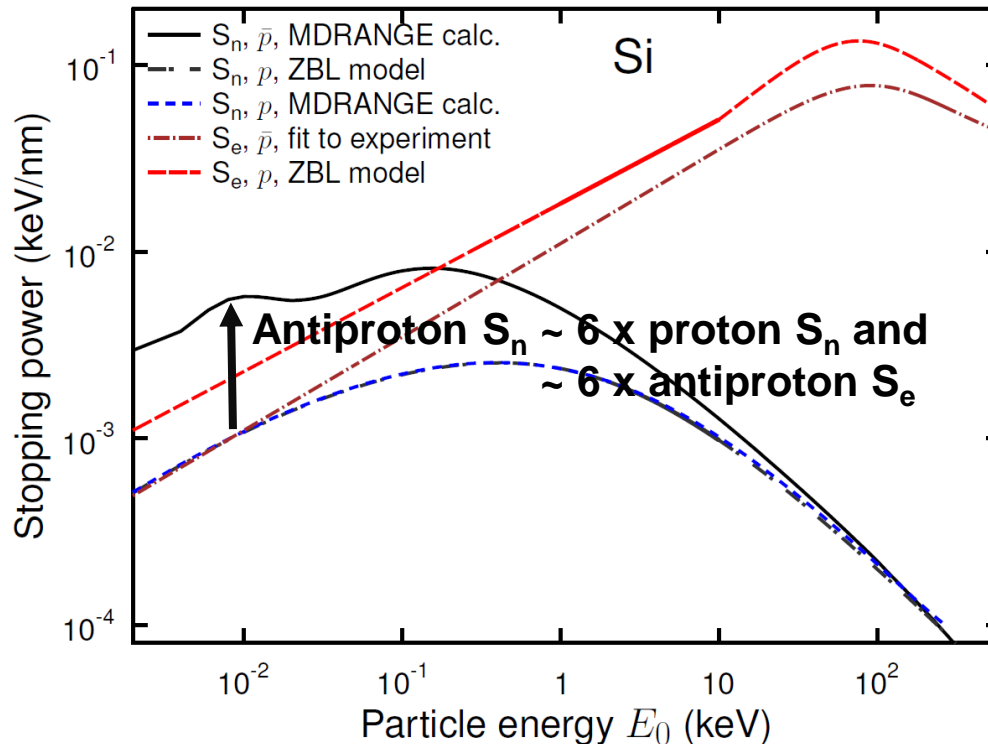


Nuclear stopping power of antiprotons: results



➤ Results show that:

- 1. The nuclear stopping power S_n of antiprotons can be stronger than their electronic stopping power (contrary to protons)
- 2. It is much stronger than that of protons



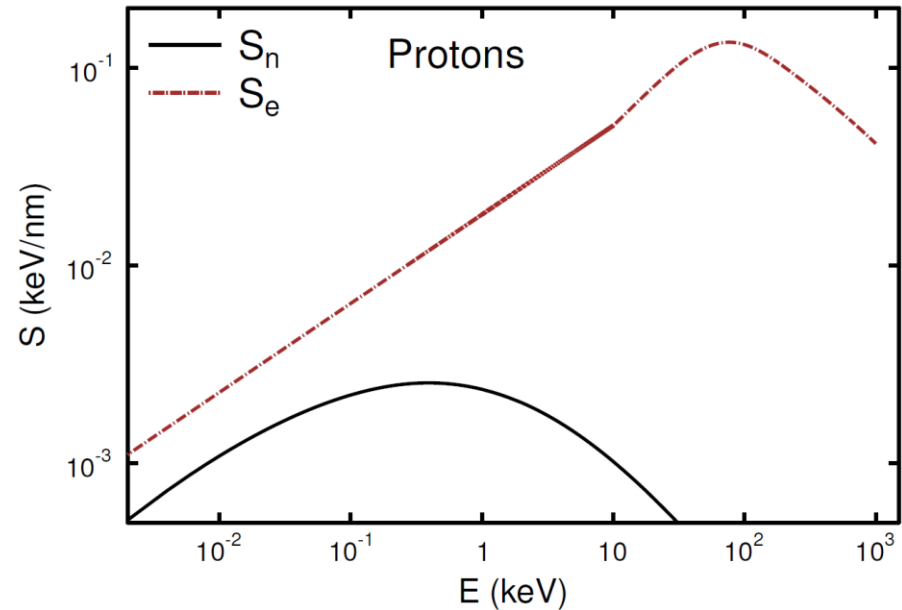
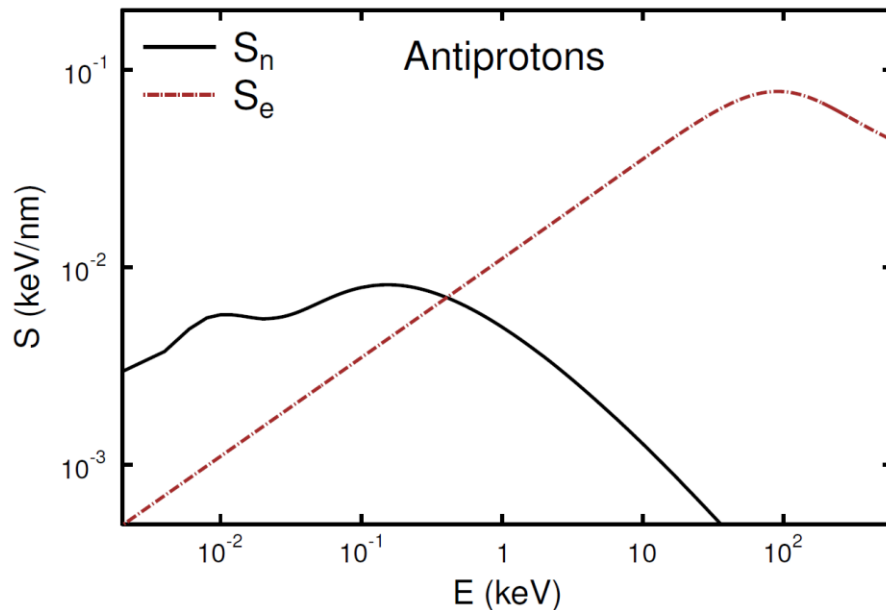
[K. Nordlund, D. Sundholm, P. Pyykkö, D. Martinez Zambrano, and F. Djurabekova, Phys. Rev. A **96**, 042717 (2017)]



Nuclear stopping power of antiprotons: results



- Summary comparison of antiprotons and protons

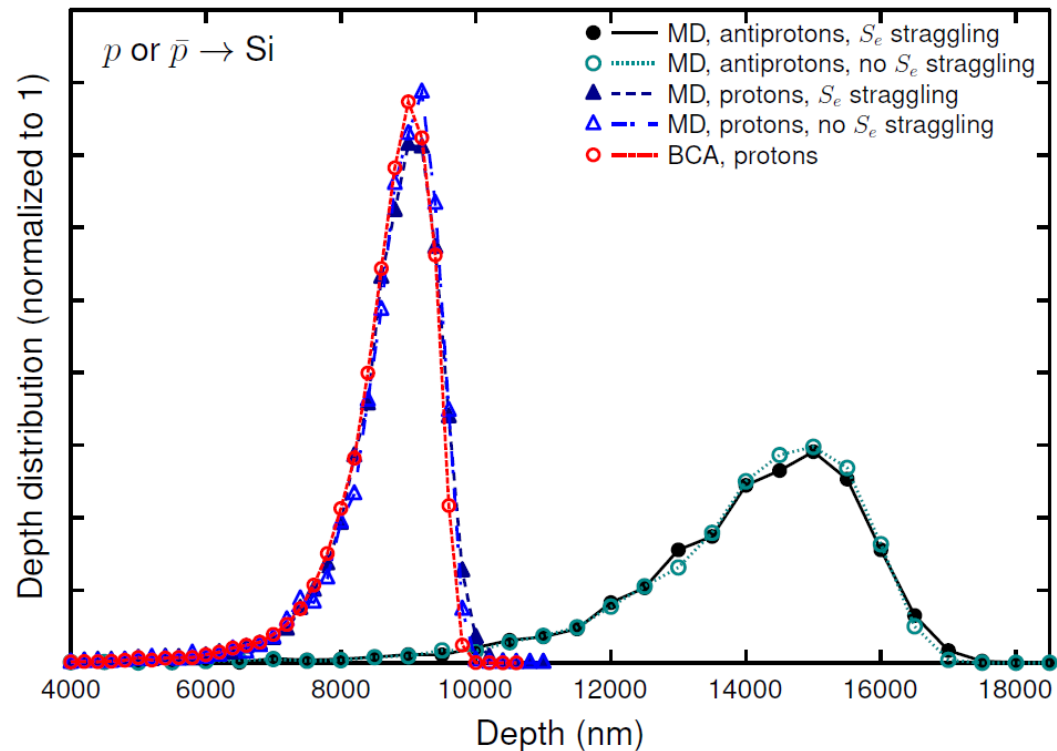




Comparison of proton and antiprotons range profiles

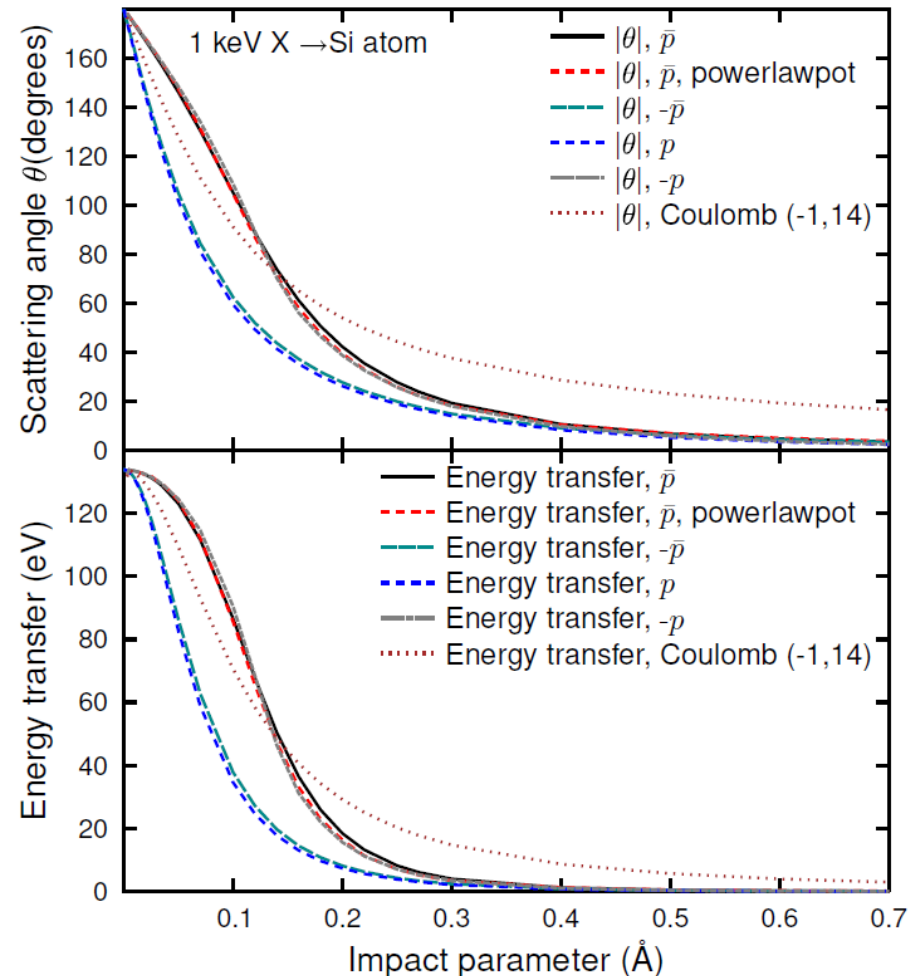


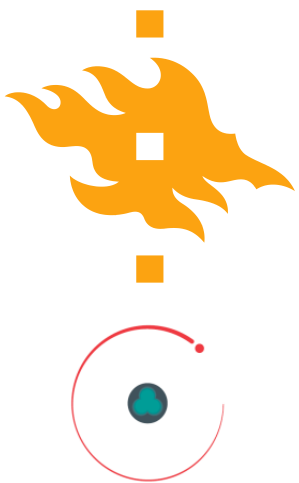
- Range profile simulations for protons give very different results, as expected



Reason to stronger stopping

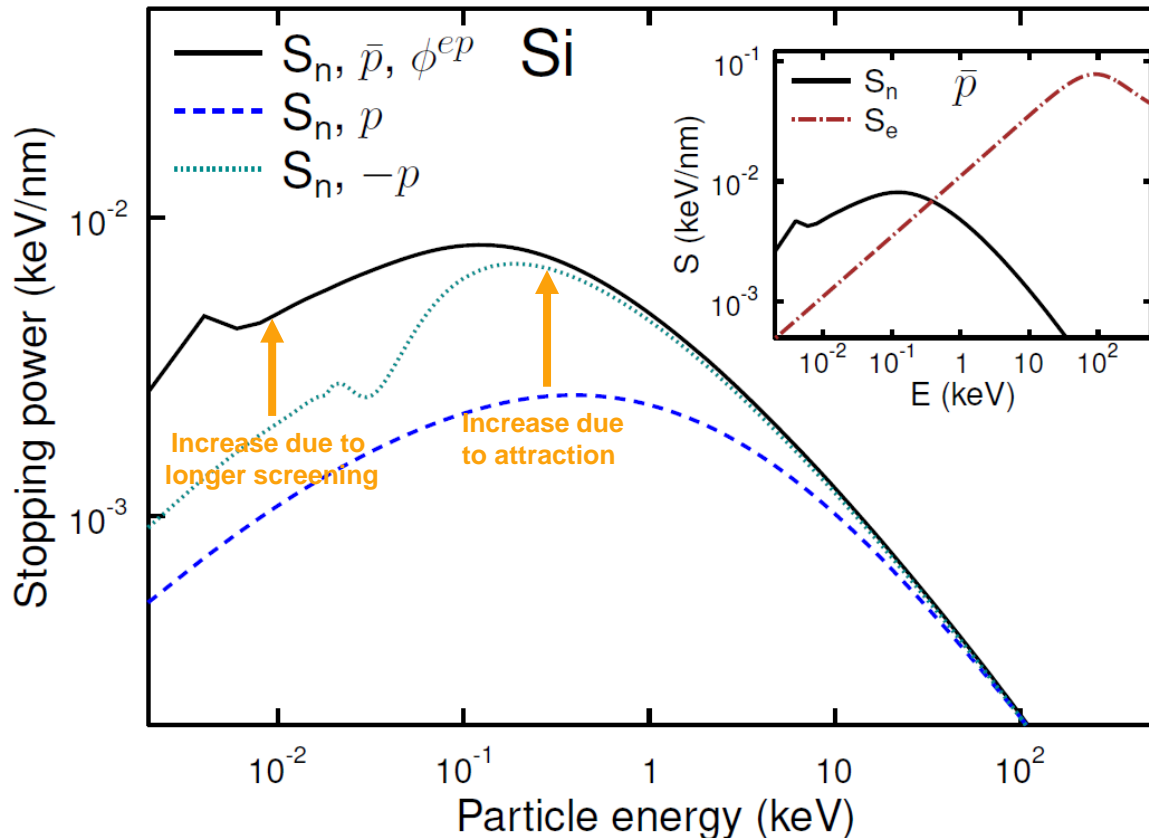
- To understand the origin of the stronger stopping, we analyzed the scattering angle and energy transfer as a function of the impact parameter for 1 keV
- Clearly larger scattering and energy transfer for antiprotons \bar{p} than protons p
- Note that a hypothetical “negative proton $-p$ ” has similar behavior as the antiproton \bar{p}





Reason to stronger stopping

- Comparison of antiprotons, protons and hypothetical “negative proton” shows that both attraction and longer screening length leads to increased nuclear stopping power





Molecular dynamics of antiproton transmission through foils



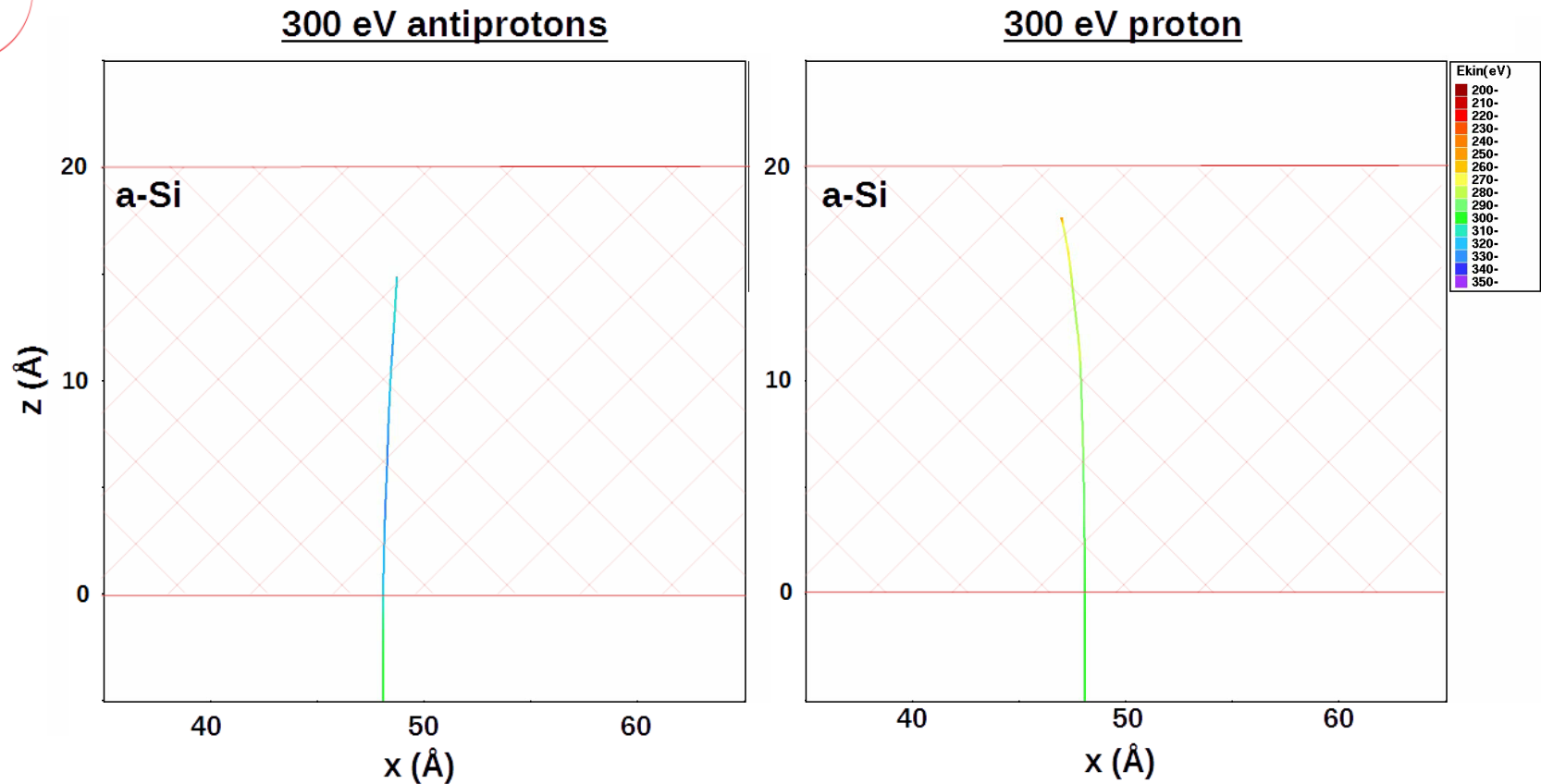
- However, to calculate the transmission through foils, one needs actual trajectory simulations, as the antiprotons do not move in straight paths
- Hence we did MDRANGE (MD in recoil interaction approximation) simulations of antiproton movement in the foils
 - In some cases, the antiprotons were captured in the purely attractive potential: observed spiral path inwards, or Kepler-like bound orbit
 - Hence cutoff criterion introduced: if antiproton came within 3 nm of nucleus, a nuclear reaction was assumed to occur and the simulation was stopped

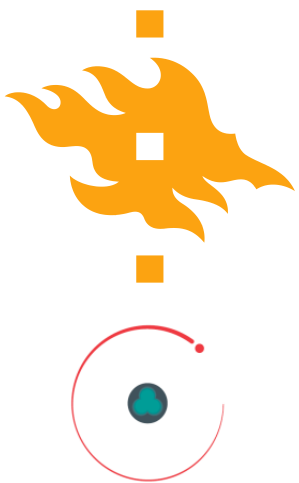


Molecular dynamics of antiproton transmission through foils

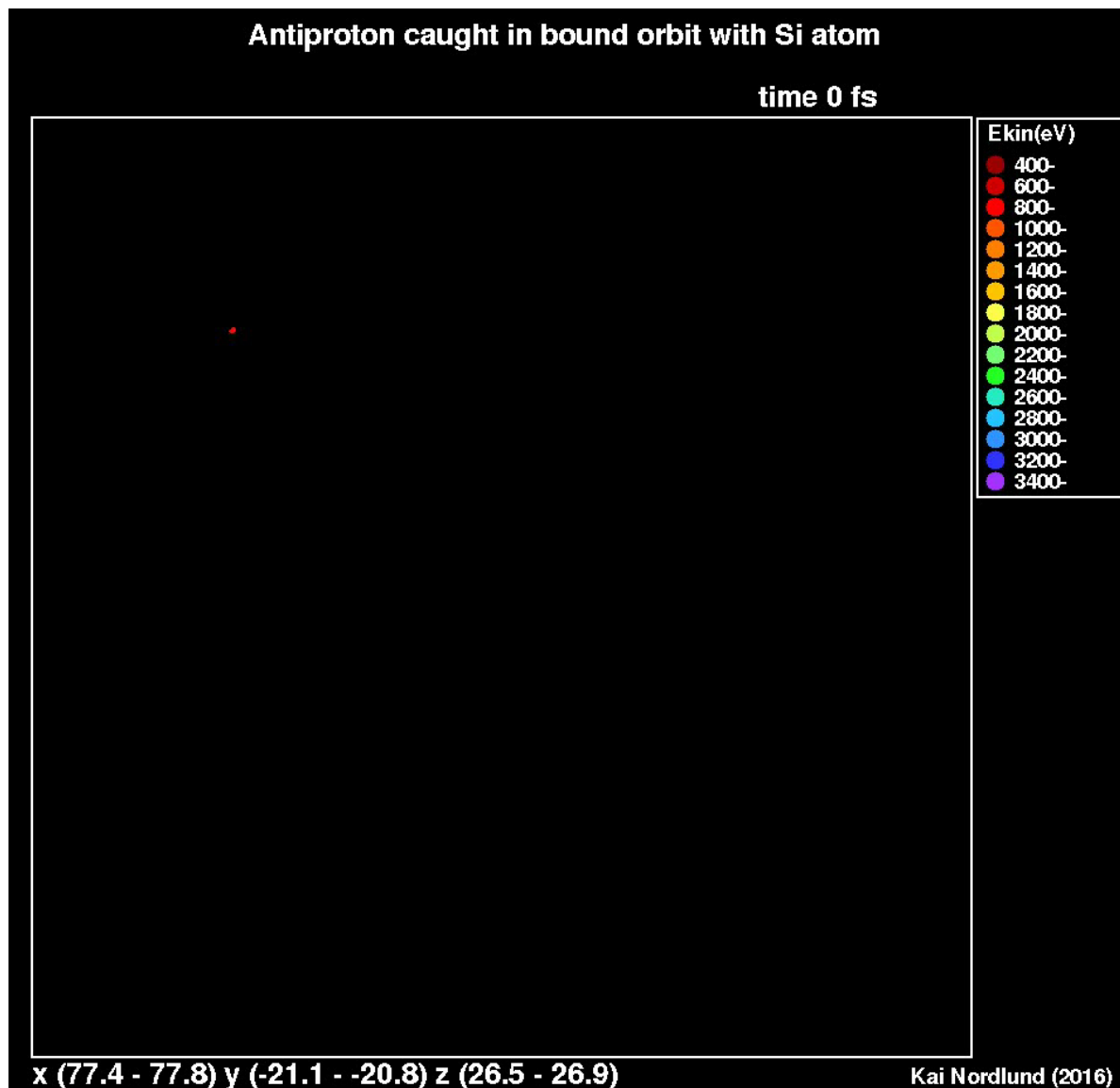


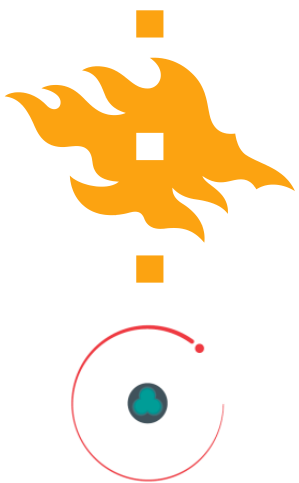
➤ Animation: transmission of 300 eV \bar{p} through 2 nm Si foil





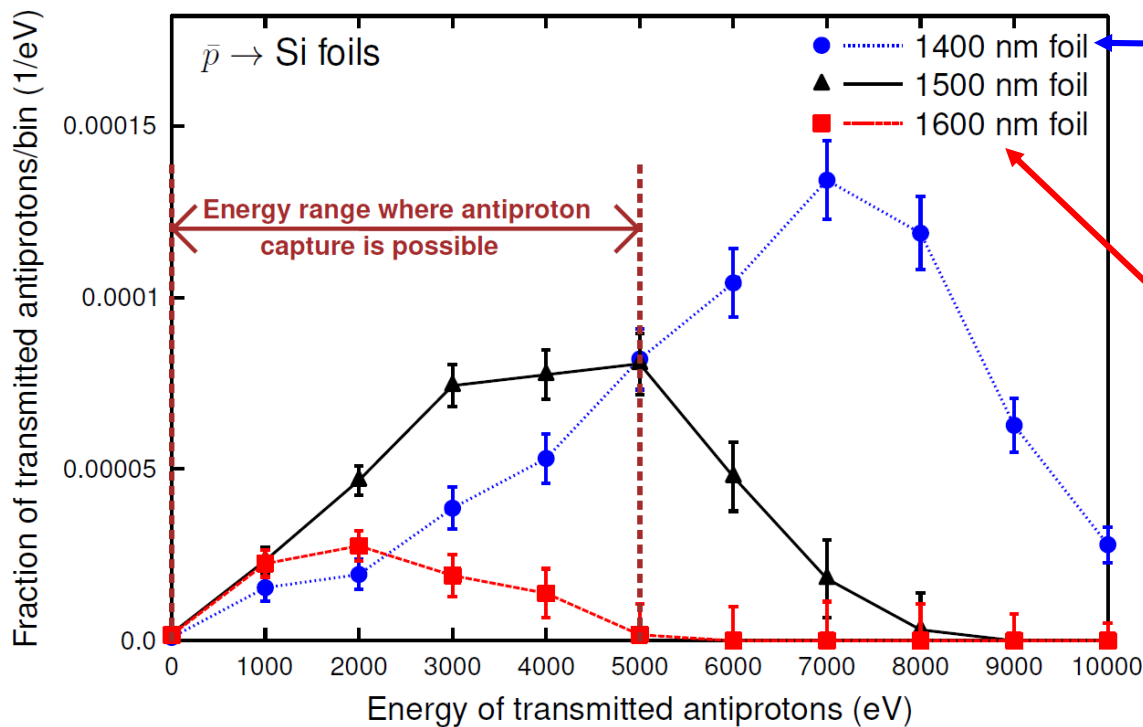
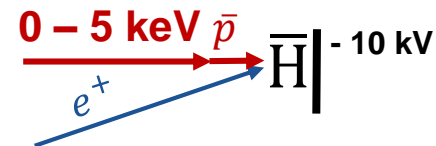
Antiproton in Kepler orbit: animation





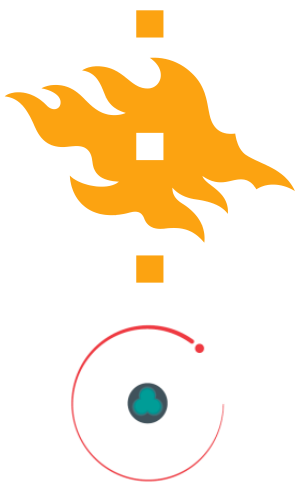
Simulations of practical importance: optimization of foil thickness, 1.

- The aim for the ATRAP experiment is to find foil materials and thicknesses that optimize for incoming 100 keV \bar{p} the transmitted fraction of 0 – 5 keV ones



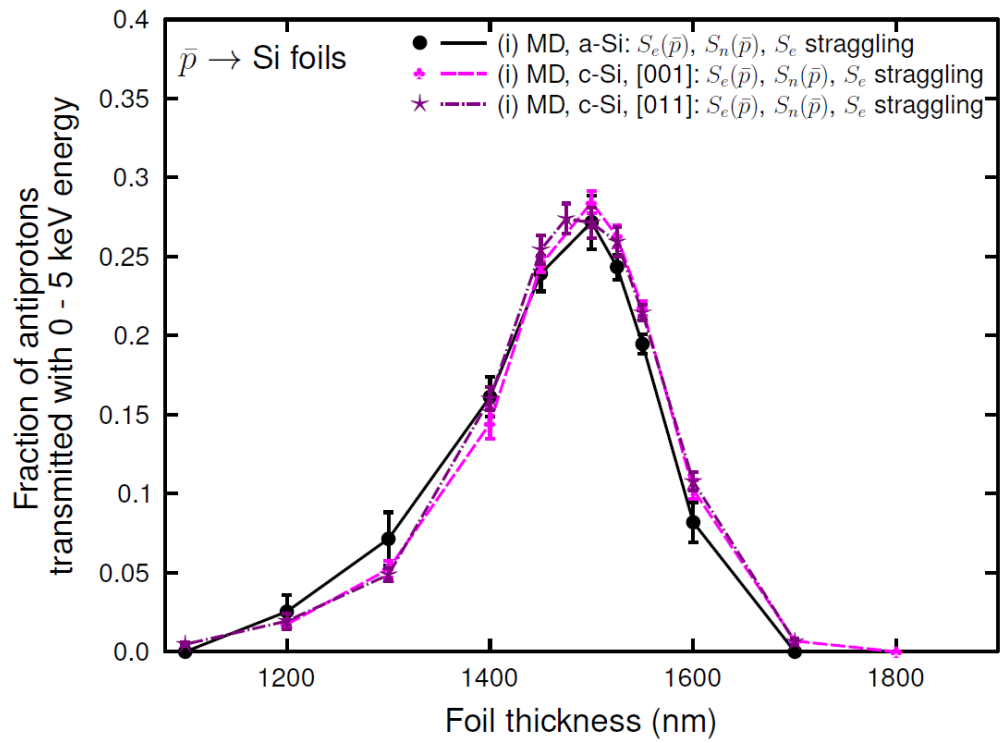
Too thin foil: energy of transmitted particles too high

Too thick foil: little or no transmission



Simulations of practical importance: optimization of foil thickness, 2.

- By integrating the fraction of antiprotons transmitted through the foil with energy between 0 and 5 keV for various thicknesses, we could find the optimal thickness
- Predicted optimal thickness for a-Si and c-Si foils: 1500 nm

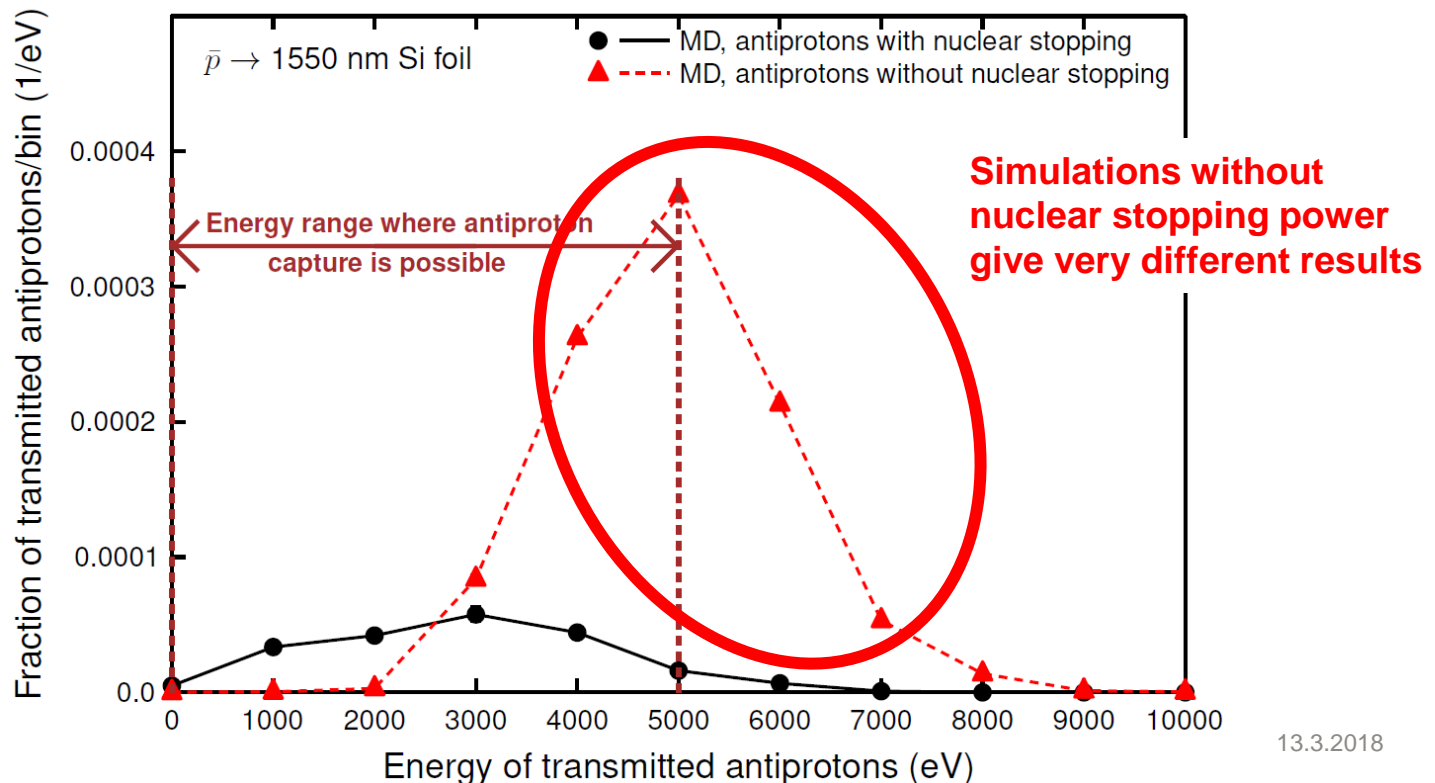


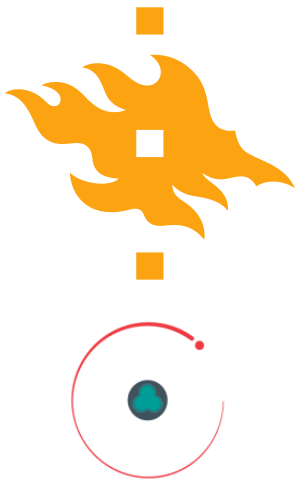


Simulations of practical importance: does the nuclear stopping actually matter?



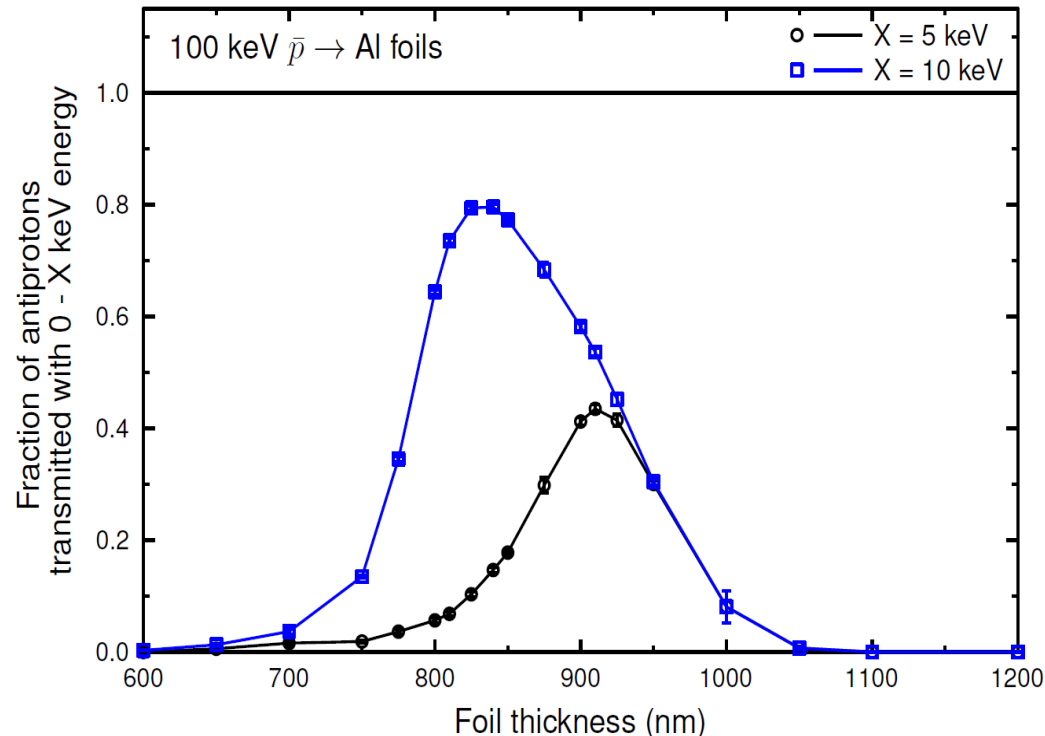
- We also tested whether doing all the trouble of including the nuclear stopping actually matters
- Answer: yes it does!

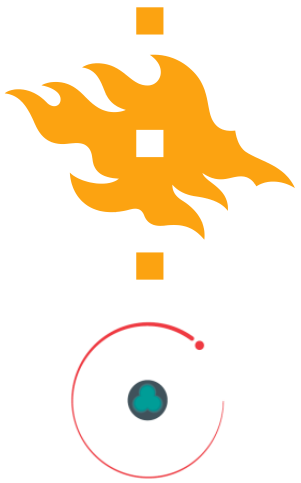




Optimization of Al

- We also did a similar optimization run for Al, as Al foils are likely easier to use than Si.
- Result: optimal thickness 900 nm (5 keV max)





Conclusions

- Using modern quantum chemistry, we determined interatomic potentials between antiprotons and 9 different elements
- These can be used for a) determining the nuclear stopping power and b) molecular dynamics simulations of antiproton movement in materials
- Results show that the nuclear stopping power of antiprotons is stronger than their electronic stopping at low energies, and also always stronger than that of protons
- Enables optimization of energy degrading foil thicknesses w.r.t. the transmitted fraction of antiprotons

[K. Nordlund, D. Sundholm, P. Pyykkö, D. Martinez Zambrano, and F. Djurabekova, Phys. Rev. A **96**, 042717 (2017)]



Backup slides





Atomic polarizability

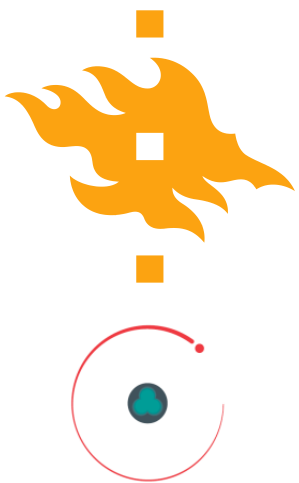


- The TurboMole calculations allowed calculating atomic polarizabilities comparable to experiments

TABLE I. Comparison of the atomic polarizabilities, α , (in a.u.) from the present Eq. (1) using MP2 data to literature values. 'PW' = present work.

| Atom | Polarizability, α | |
|------|--------------------------|-------------------------|
| | PW | Lit. |
| Be | 40.90 | 37.76 ^a |
| N | 5.34 | 7.63 ^b |
| Ne | 2.44 | 2.66110(1) ^c |

^a Ref. [41], ^b Ref. [42] (exp), ^c Ref. [43] (exp).



Atomic polarizability

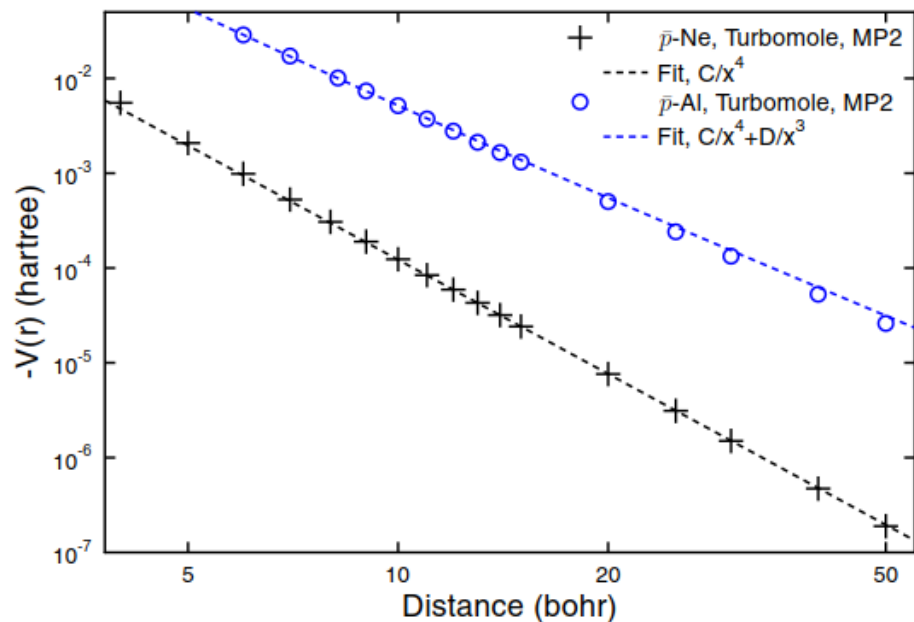
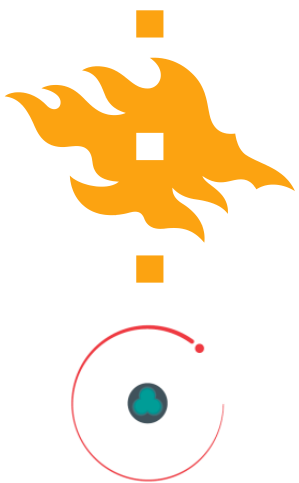


FIG. 4. The calculated MP2 antiproton-neon and antiproton-aluminum potentials (symbols) at large separations and a fit to $V(r) = -C/2r^4$ with $C = 1.222$ hartree for Ne and $V(r) = -C_3/r^3 - C_4/r^4$ with $C_3 = 3.611$ and $C_4 = 15.675$ hartree for Al.



Different screenings

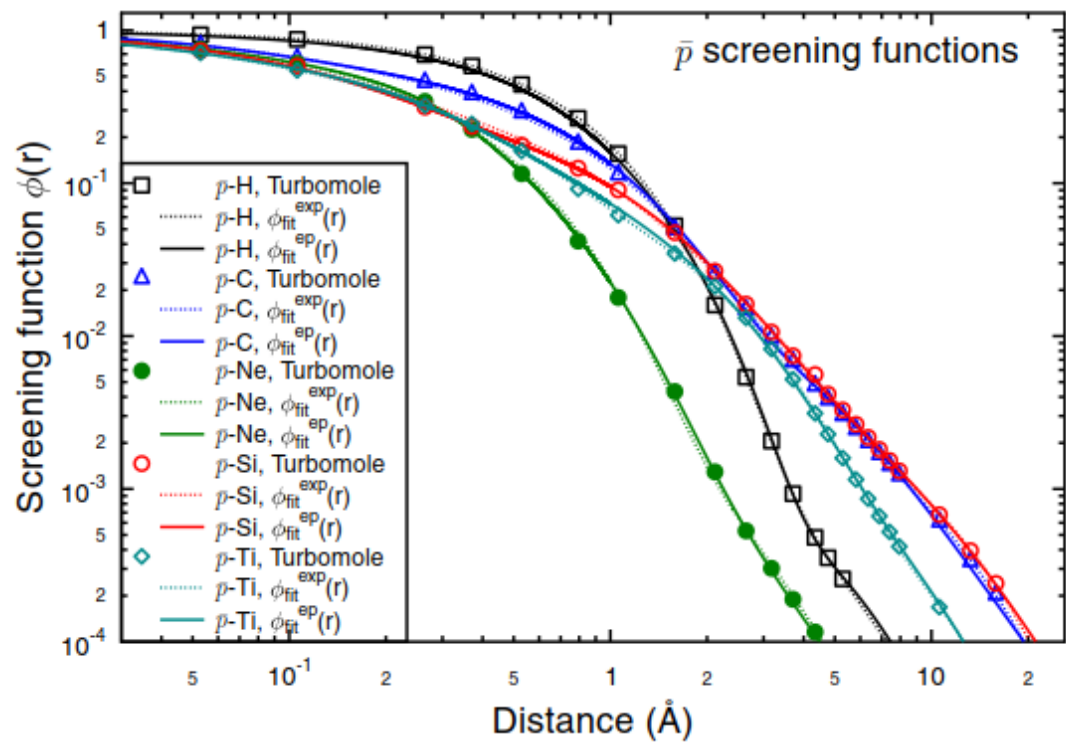
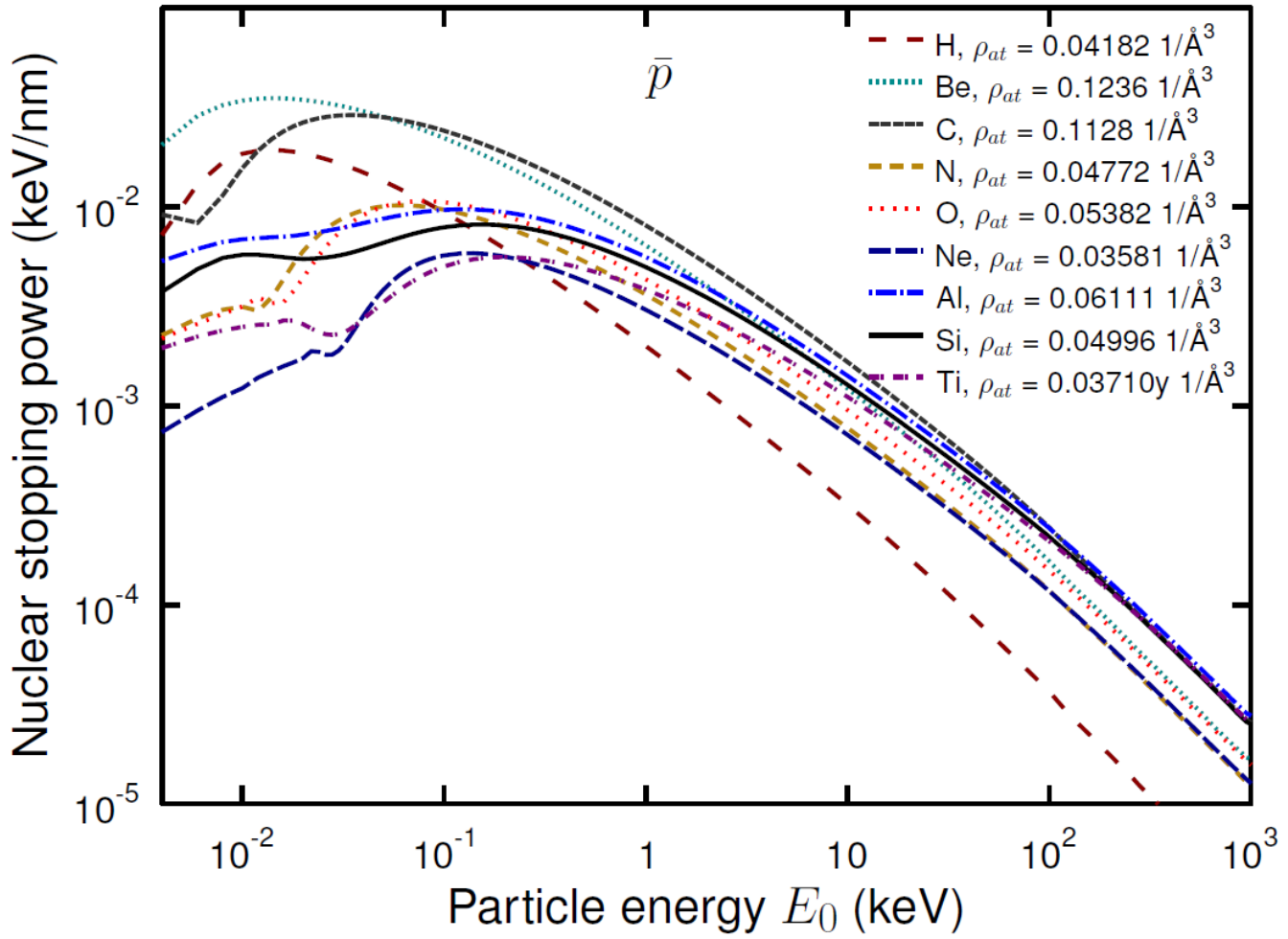


FIG. 5. Comparison of screening function fits obtained with the two different functional forms $\phi_{\text{fit}}^{\text{exp}}$ and $\phi_{\text{fit}}^{\text{ep}}$.

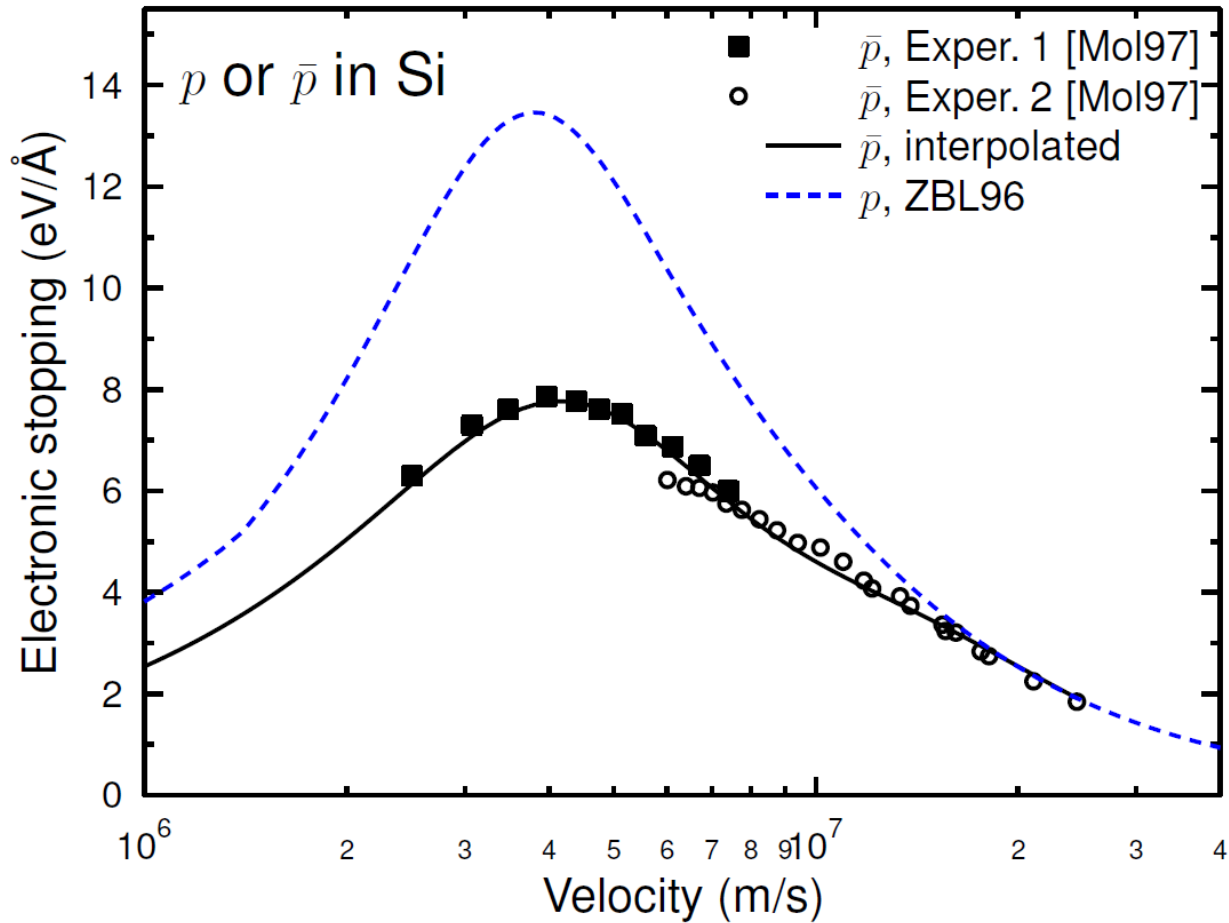


All nuclear stopping powers





Electronic stopping of antiprotons in Si

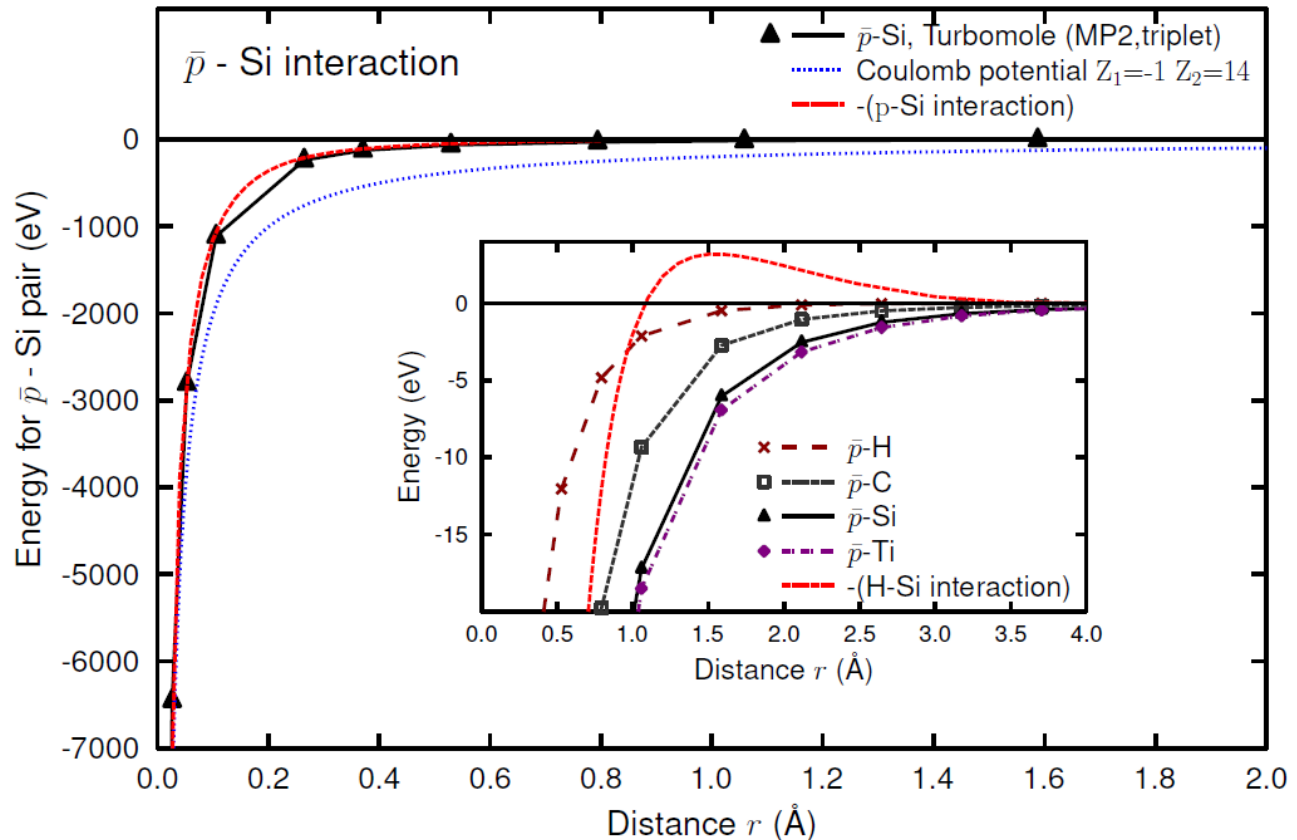




Comparison with p-Si and H-Si

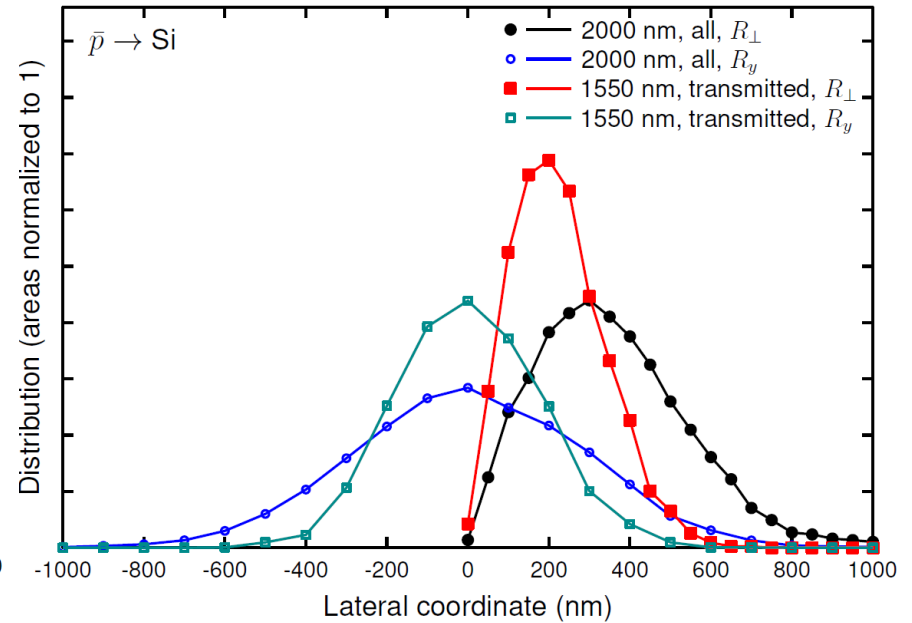
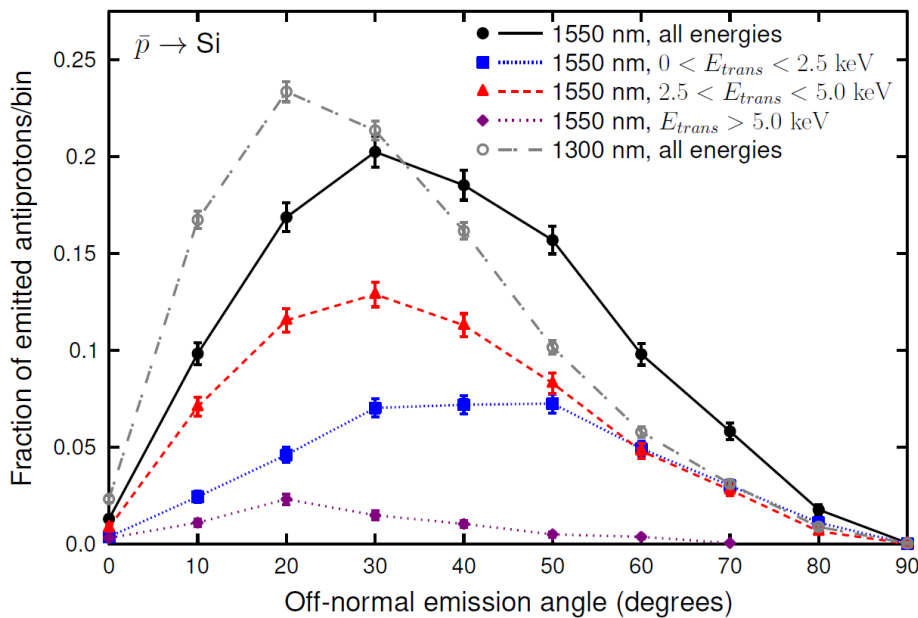


➤ Comparison with p-Si and H-Si



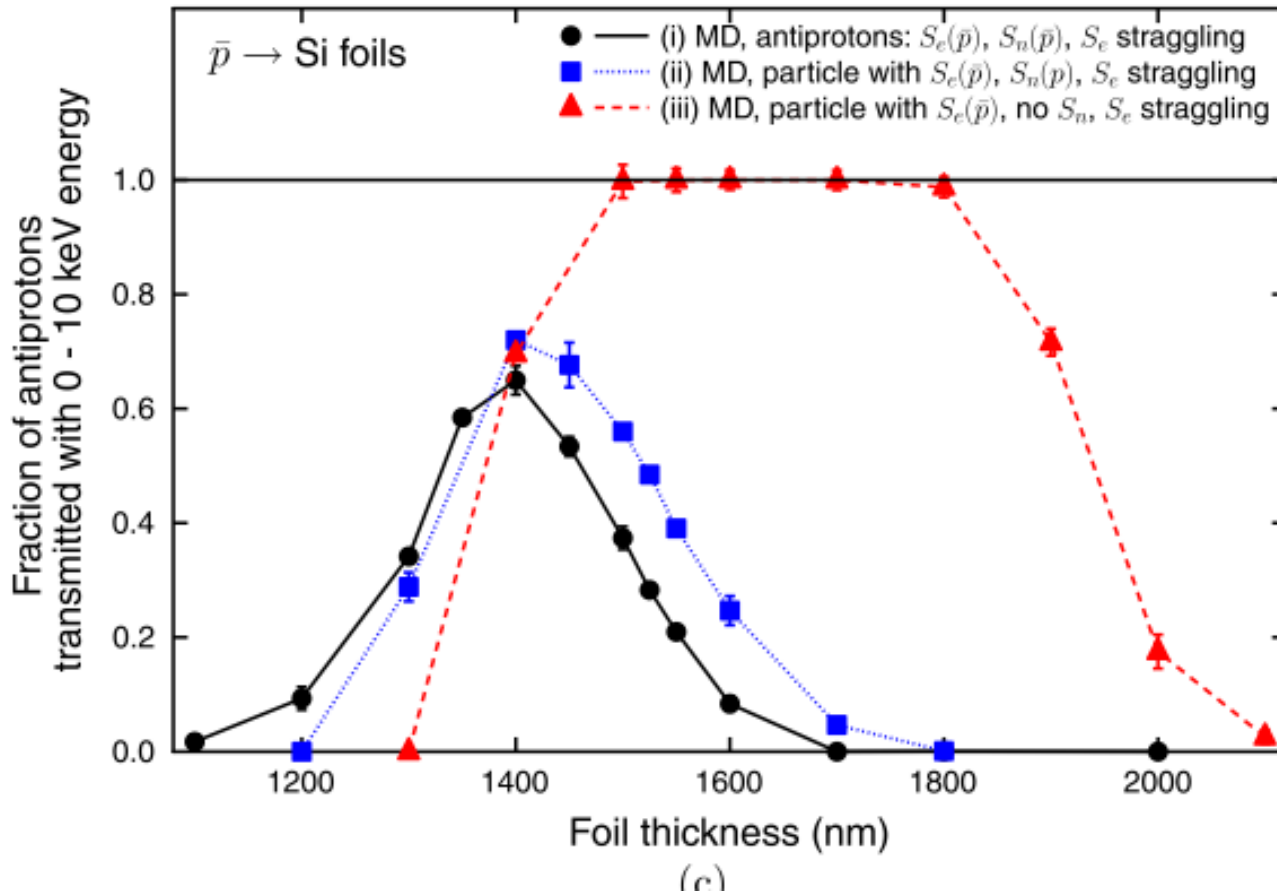


Angular and lateral spread on exit





With 10 keV upper limit





Background: antihydrogen production at CERN

