

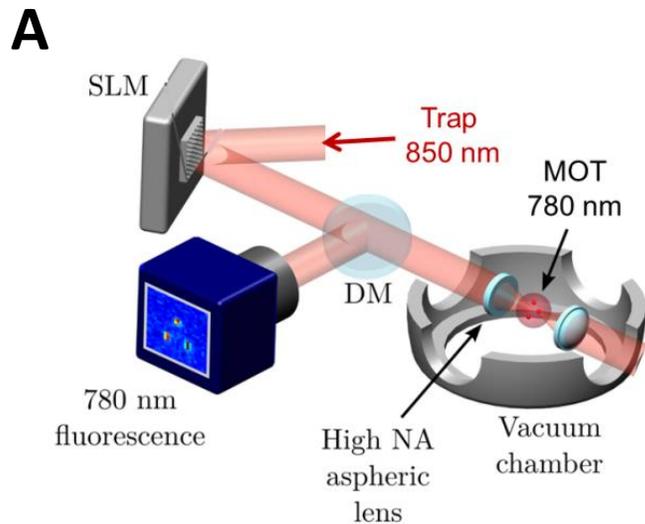
Towards Arrays of Cryogenic Traps for Improved Quantum Simulation

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Arrays of single atoms



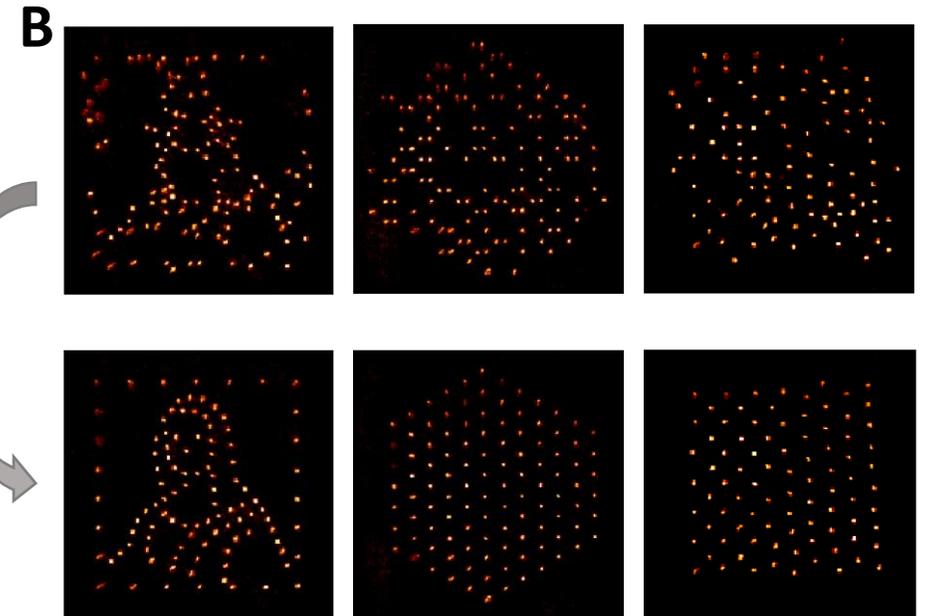
- *Up to 200 atoms, spacing $\sim 10 \mu\text{m}$*

- *Interactions: Rydberg states*

- *Atom-by-atom sorting into ordered arrays with a moving tweezers:*

- *Extension to 3D arrays*

- *Now: cryogenic platform*



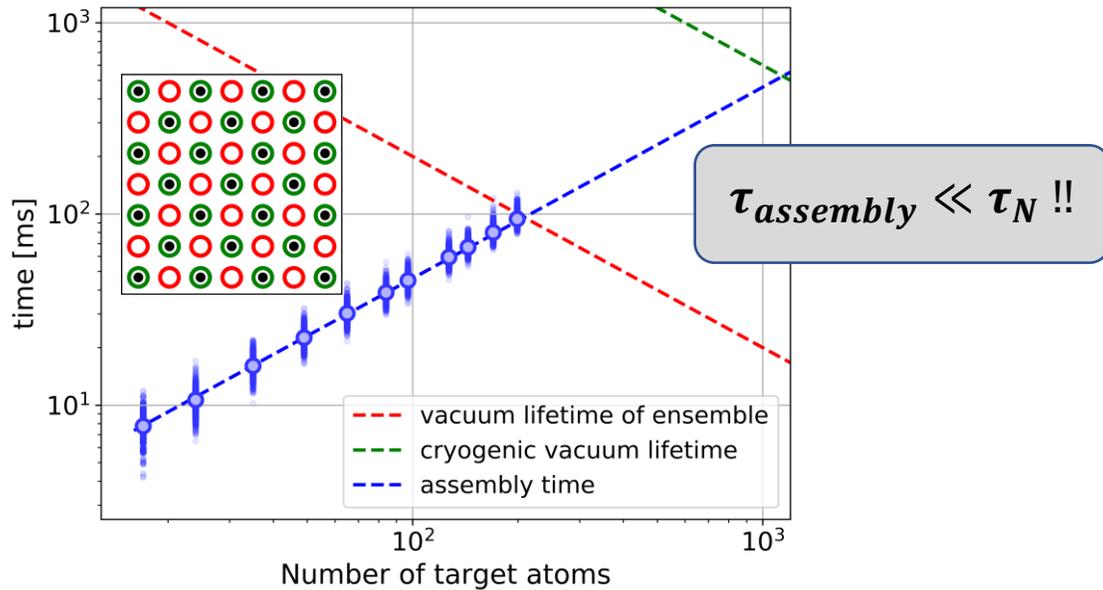
D. Barredo *et al.*, *Science* 2016

K. Schymik *et al.*, In preparation (2020)

Why Cryogenic ?

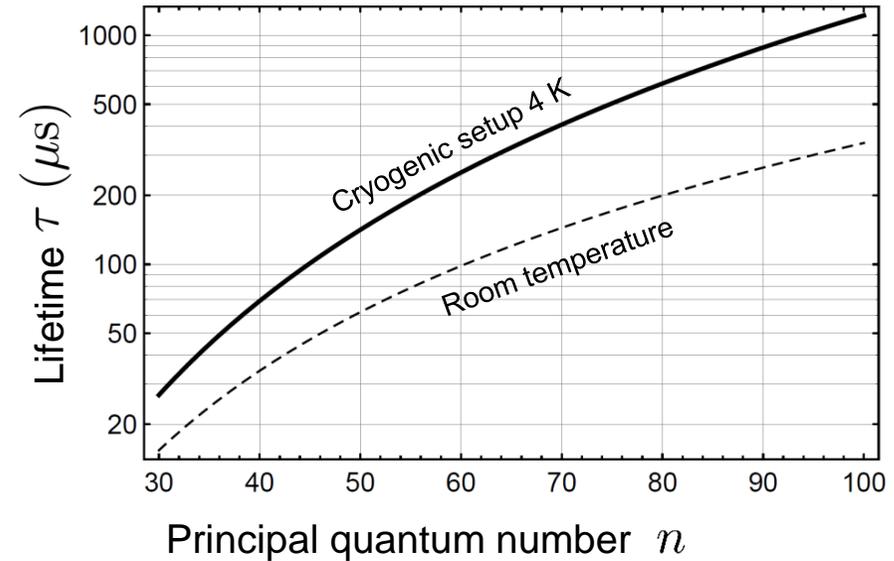
C When scaling up the number of atoms N , the single-atom vacuum-limited lifetime τ is a limitation!

- Current room-temperature setup: $\tau \approx 20$ s
- Lifetime of N -atom array: $\tau_N = \tau/N$
- Assembly time of N -atom array: $\tau_{assembly} \propto N$



- Cryogenic lifetime $\tau_{cryo} \gg \tau$ due to better vacuum!

D Increased Rydberg lifetime

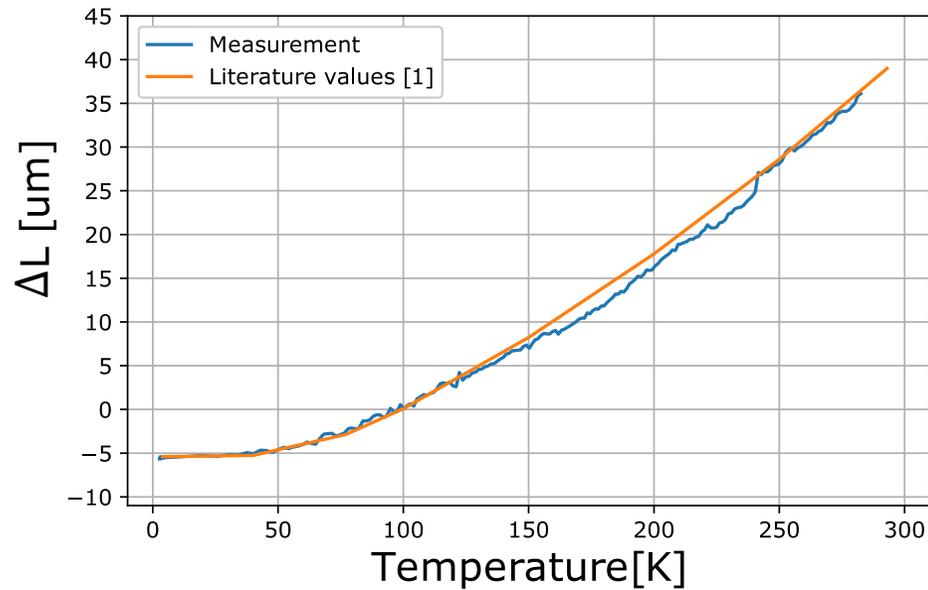
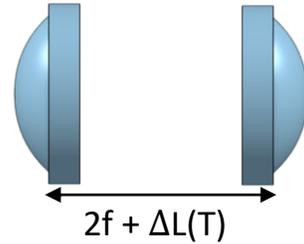


- Reduce black-body-induced transitions to neighboring states
- Increased coherence for quantum simulation

Technical Challenges

E Thermal contraction

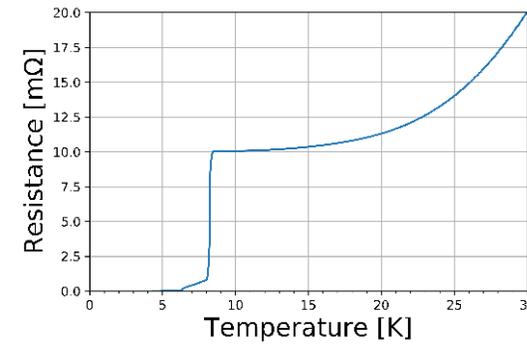
- Stress-free mount of lenses
- Compensation of 2-f configuration



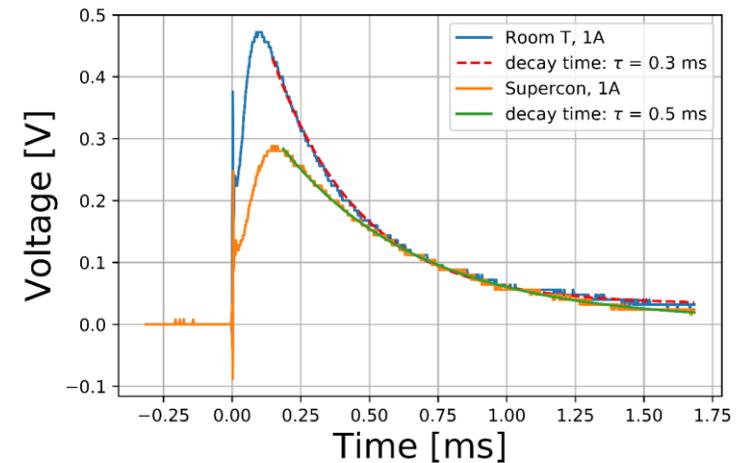
[1] R.J. Corruccini, J.J. Gniewek. *Thermal Expansion of Technical Solids at Low Temperatures*. National Bureau of Standards Monograph 29 (1961).

F Superconducting coils: No heat load!

- Transition temperature at 9 K



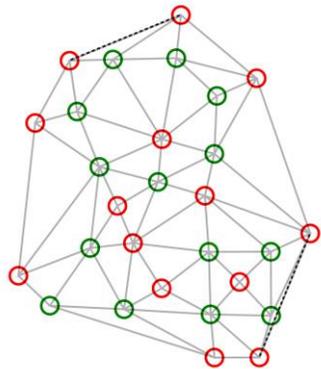
- Eddy currents decay ($\tau = 0.5$ ms)
 - measurement with pick-up coil



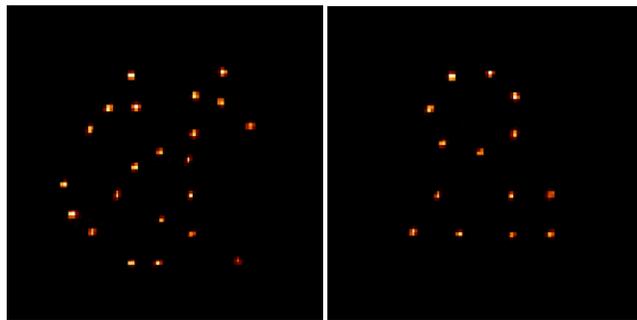
Improved Assembler

G Arbitrary patterns due to graph approach

- **Change from Bravais lattice based algorithms**
- **Interesting e.g. for:**
 - crystal defects (dislocation, grain boundary, ...)
 - disordered Arrays (e.g. Anderson localization studies)
 - Completely arbitrary arrays for combinatorial optimization problems (maximum independent set)



Red: reservoir traps
Green: target traps



Initial loading

Assembled Array

H Improved algorithms

- **Goal: decrease number of elementary moves**
 - Pick-up and release time dominant timescale
- **Best algorithm depends also on target structure**

