

HYSPPLIT Implementation and Verification for HazDAC and SUMMIT

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Overview

- Current rising junior at Princeton University
 - Majoring in Chemical Engineering
 - Minors in computer science and materials science
- Working on incorporating models into toolkits
 - HazDAC
 - SUMMIT
- Verifying that the models work correctly and are consistent
 - HYSPLIT
 - HPAC

Bioterrorism Preparation

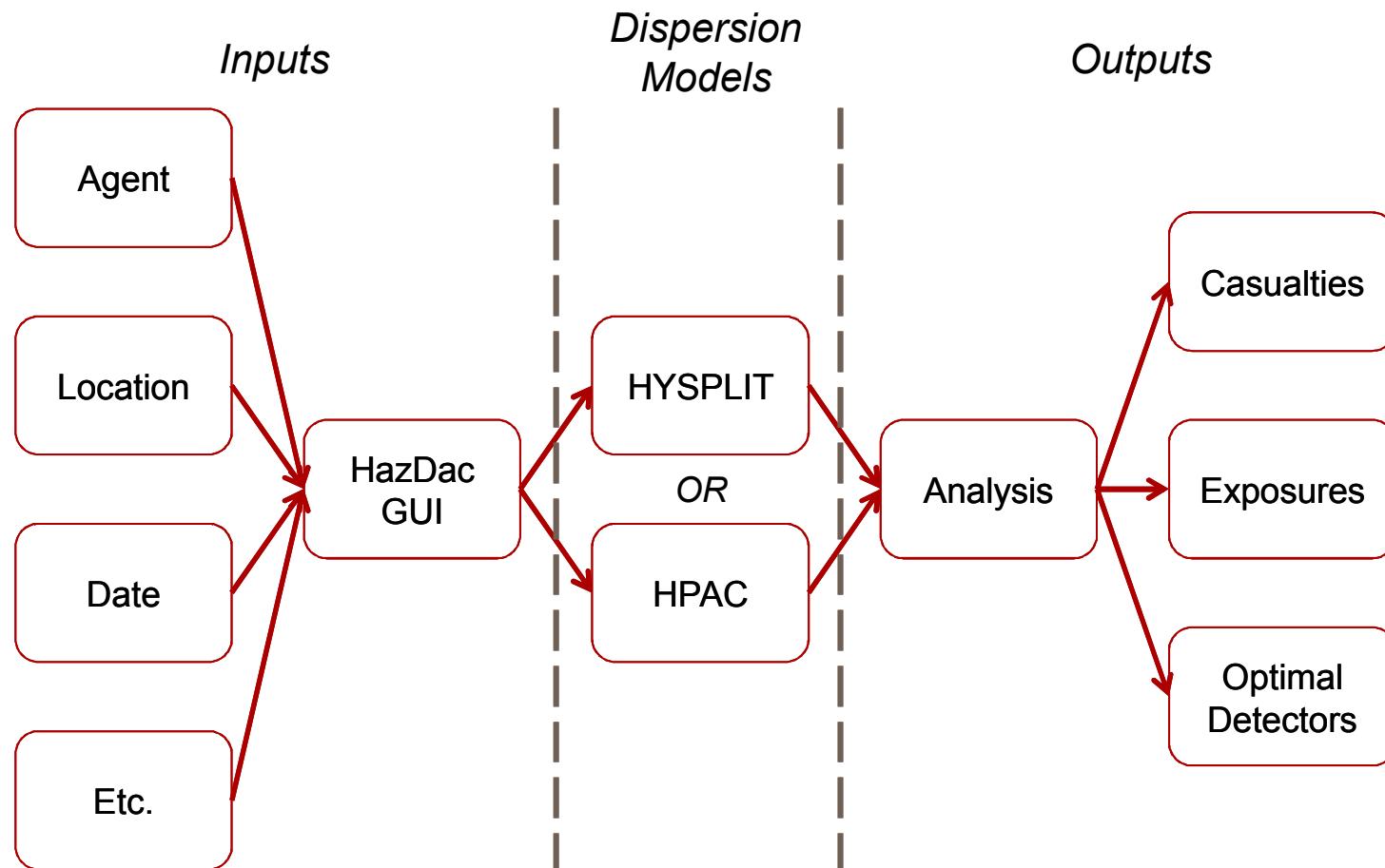
- Need to be prepared for possible agent release
 - Where to place sensors to best detect a release?
 - How should emergency response teams prepare?
- Two Options:
 - Tracer experiments
 - Expensive and time intensive
 - Only accounts for one place/time
 - But is real
 - Simulations
 - Can account for any place/time
 - Cheap and not time intensive
 - But only as good as algorithm and input data

Modeling as a Preparation Tool

- Many models simulate the release of an agent
- HPAC – various agent release simulations
 - DTRA created
 - Standard for threat modelling
 - But a pain to install, only works on 32-bit windows
- HYSPLIT – atmospheric dispersion simulations
 - NOAA created
 - Platform independent and very lightweight

HazDAC Overview

- Integrated toolset for simulating chem and bio incidents and responses



Verification of HYSPLIT Necessary

- HPAC
 - HazDAC default model
 - Assume accurate
- *NEW* HYSPLIT
 - Much easier to install for clients
- Can we trust HYSPLIT?
 - Need to affirm that HYSPLIT is “equivalent” to HPAC for HazDAC’s needs

Visualization of HPAC and HYSPLIT

Location: Albuquerque (ABQ)

Duration: 12.0 hrs

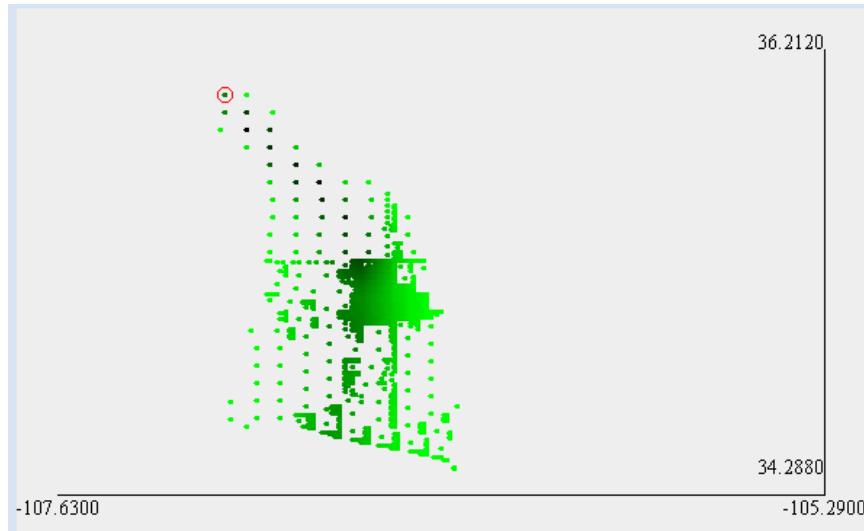
Agent: Anthrax

Start date: 01/01/08 03:00:00

Concentration normalized to a scale of 0 – 1 then translated to colored pixel.

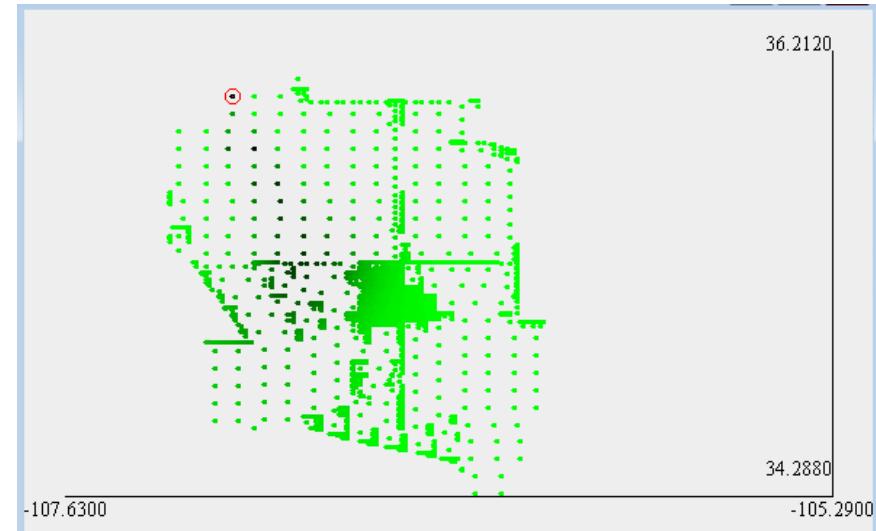
HYSPLIT:

(normalized to 4.9E-7, MAX = 4.9E-7, MIN = 3.23E-13)



HPAC:

(normalized to 8.9E-7, MAX = 8.9E-6, MIN = 1.0E-13)



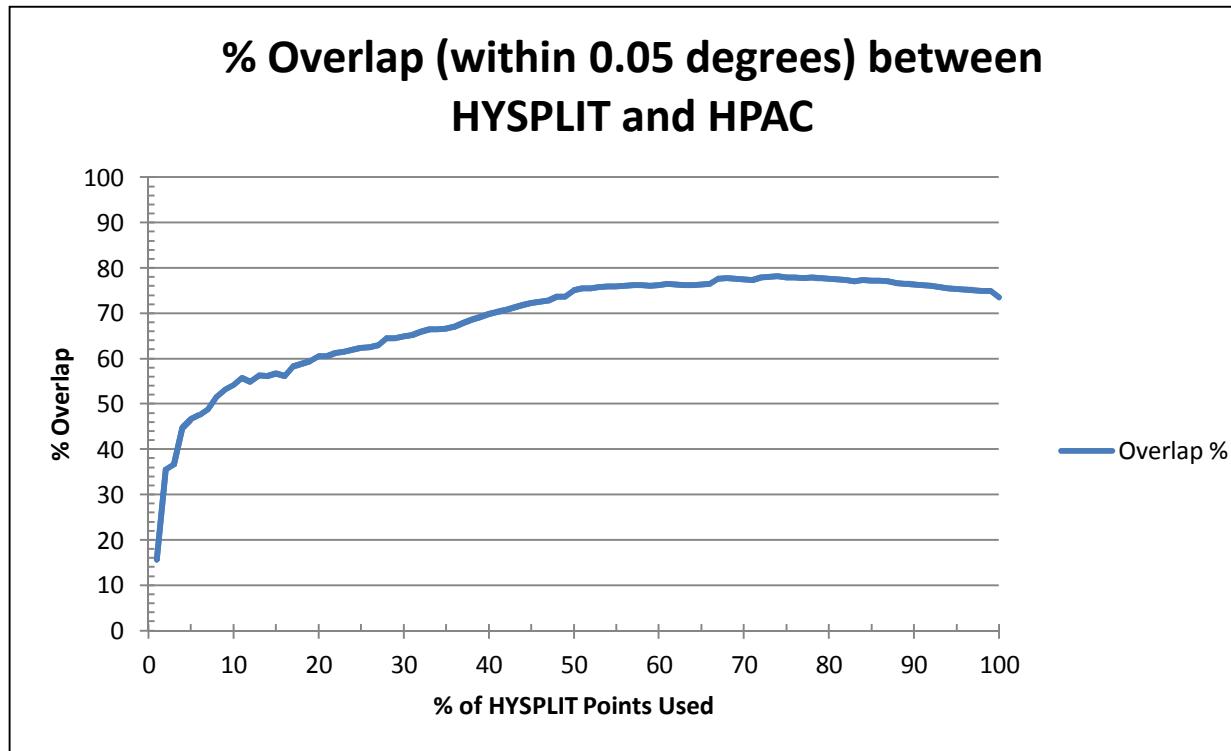
* Four points excluded from normalization in HPAC run

Quantifying Comparisons

- Concentration overlap
 - How much of the top x% points are the same in HYSPLIT and HPAC?
- Concentration profile
 - How do the absolute concentration numbers compare?
- Exposures and casualties
 - How do they compare?
- Detection optimization
 - How does the percent detected compare?
 - How do the optimal detection locations compare?
- Remember: HPAC and HYSPLIT will **never** be exactly the same

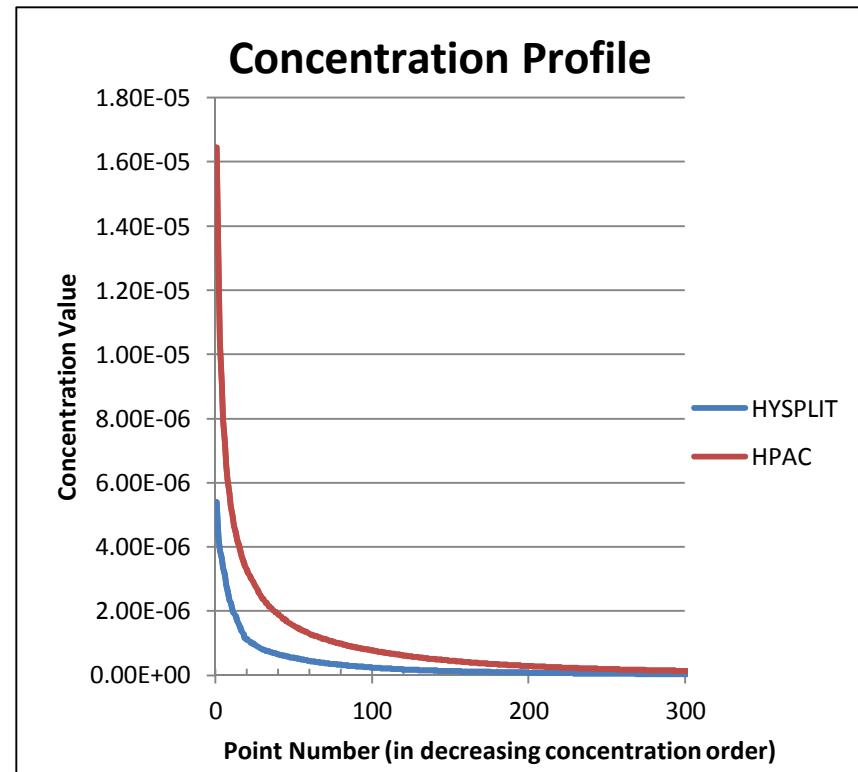
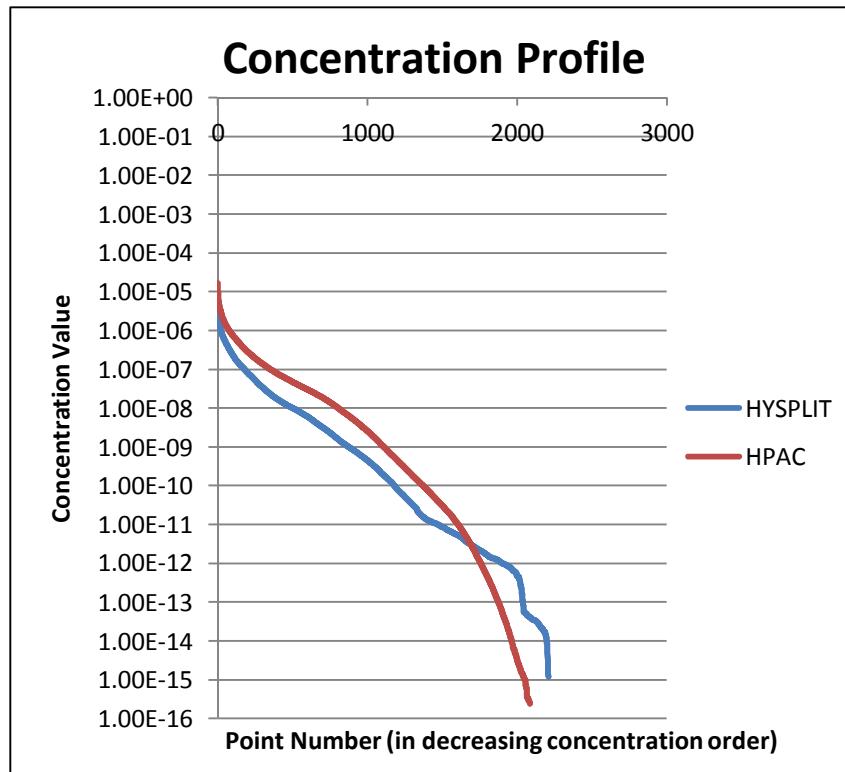
Similar Concentration Overlap

- Significant overlap in the regions covered by both models



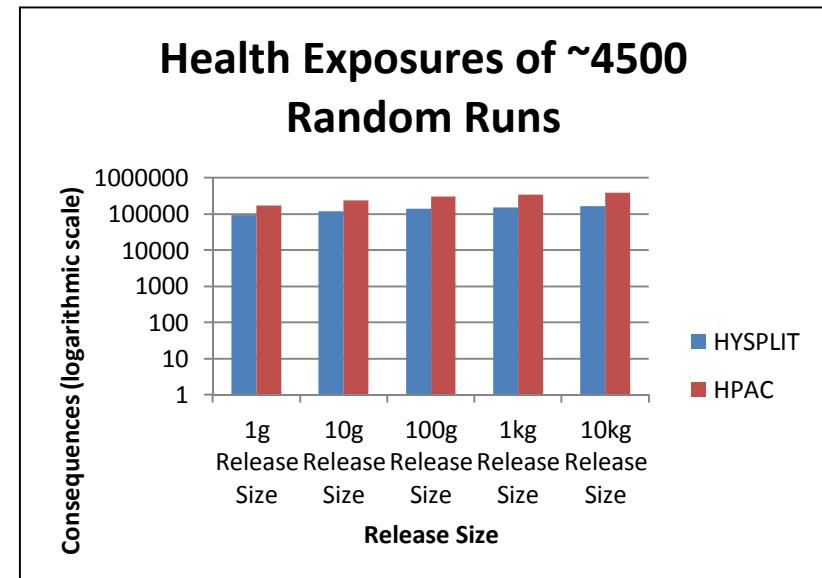
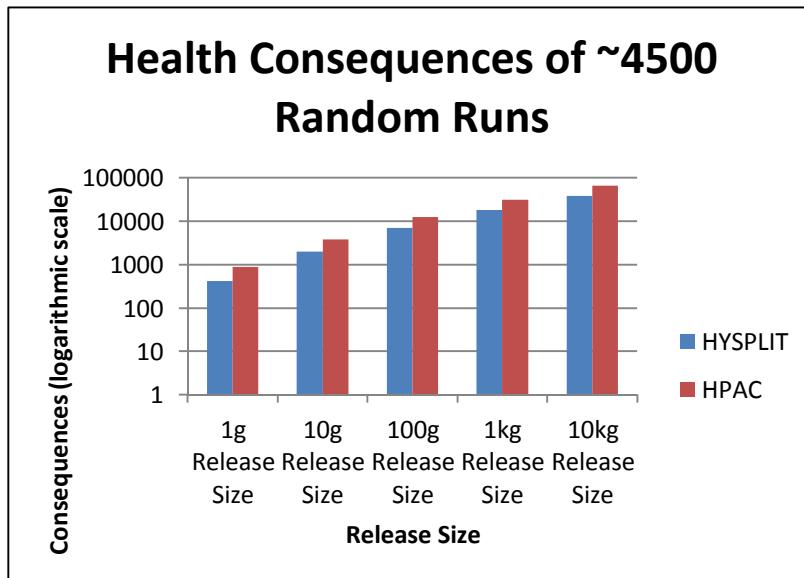
Larger HPAC Concentration Profile

- HPAC appears to have more “concentration” than HYSPLIT



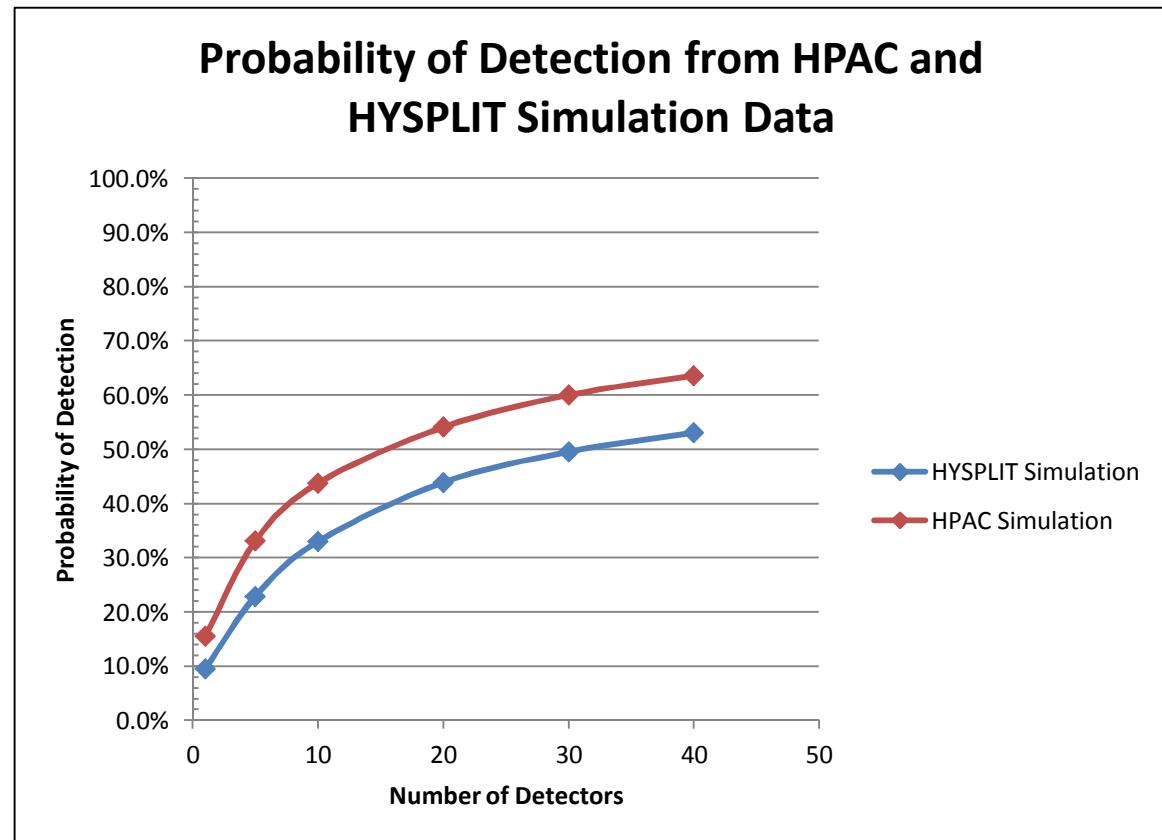
Larger HPAC Health Effects

- HYSPLIT and HPAC are same order of magnitude
- But HPAC consistently ~2 times more than HYSPLIT



Higher HPAC Detector Optimization

- A *very* good metric of comparison
- ~4500 random runs in ABQ

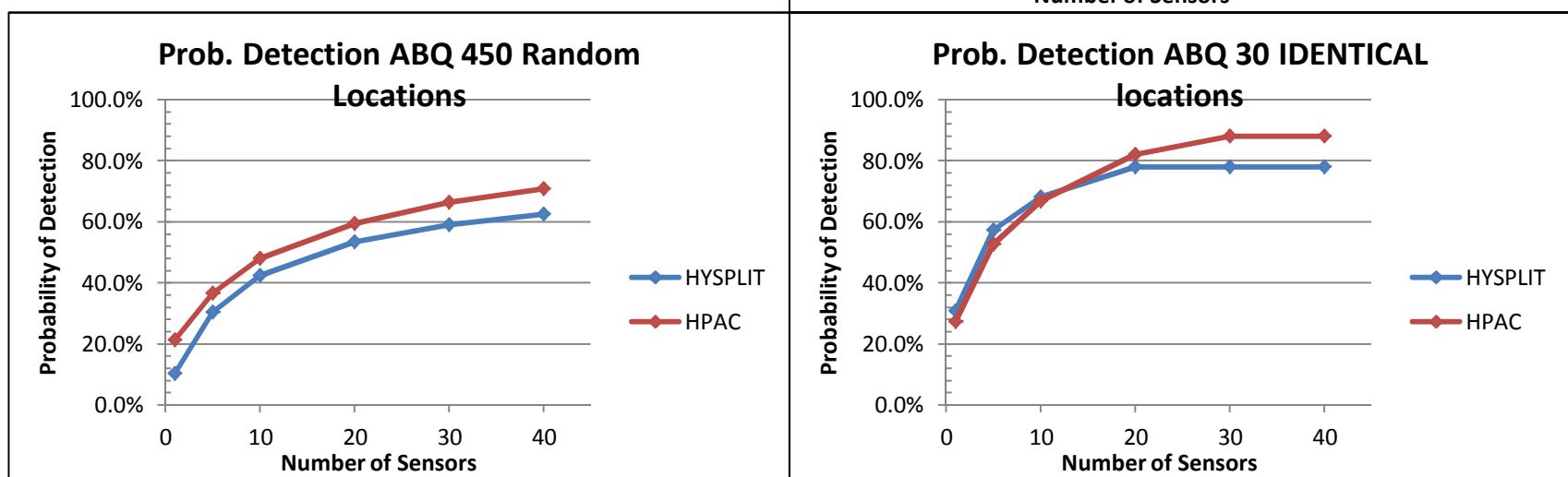
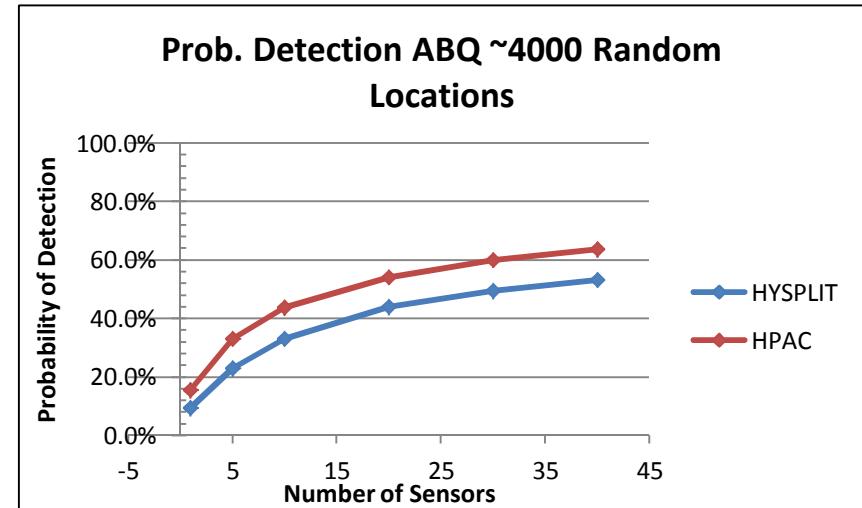


Model Differences: Consistent?

- HYSPLIT and HPAC have a reasonable overlap
- The exposure and consequence numbers are comparable
- Yet HPAC leads to higher probabilities of detection
 - Also has consistently higher exposure and consequence numbers
 - Could be caused by larger concentration values
- Next Steps:
 - Find reasonable run size
 - Check consistency of results

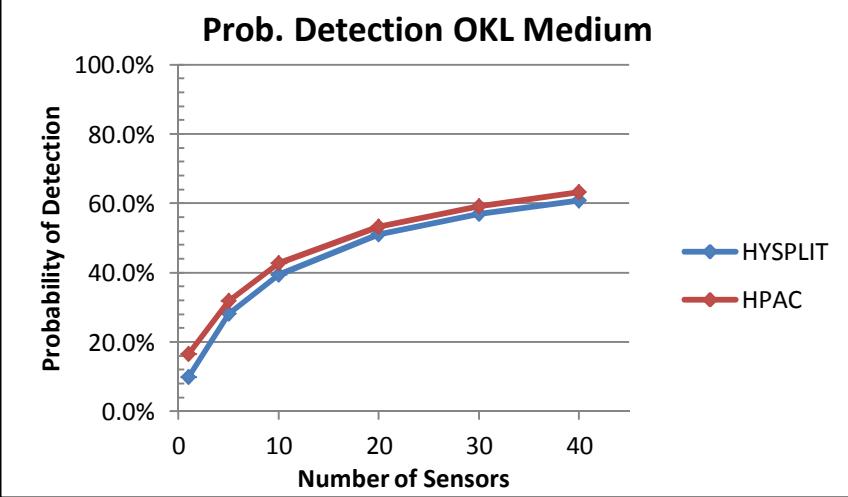
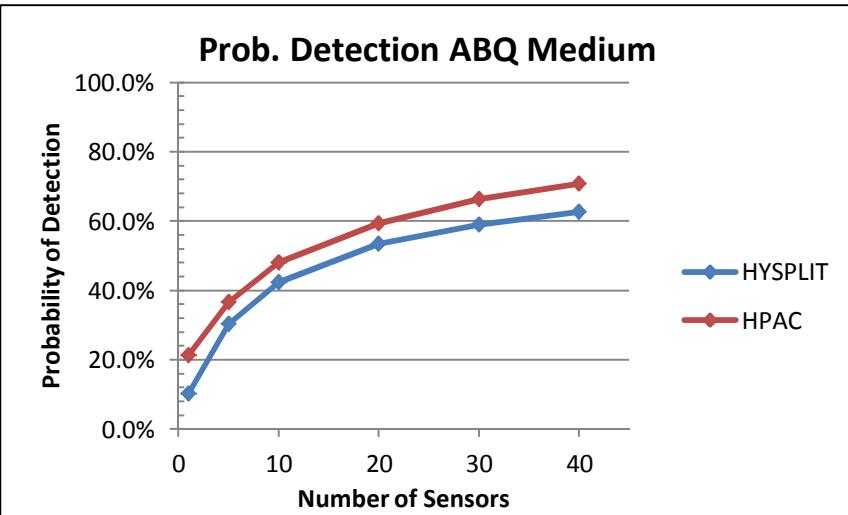
Reasonable Run Sample - 450

- Memory
- Time
- 450 is a reasonable run size



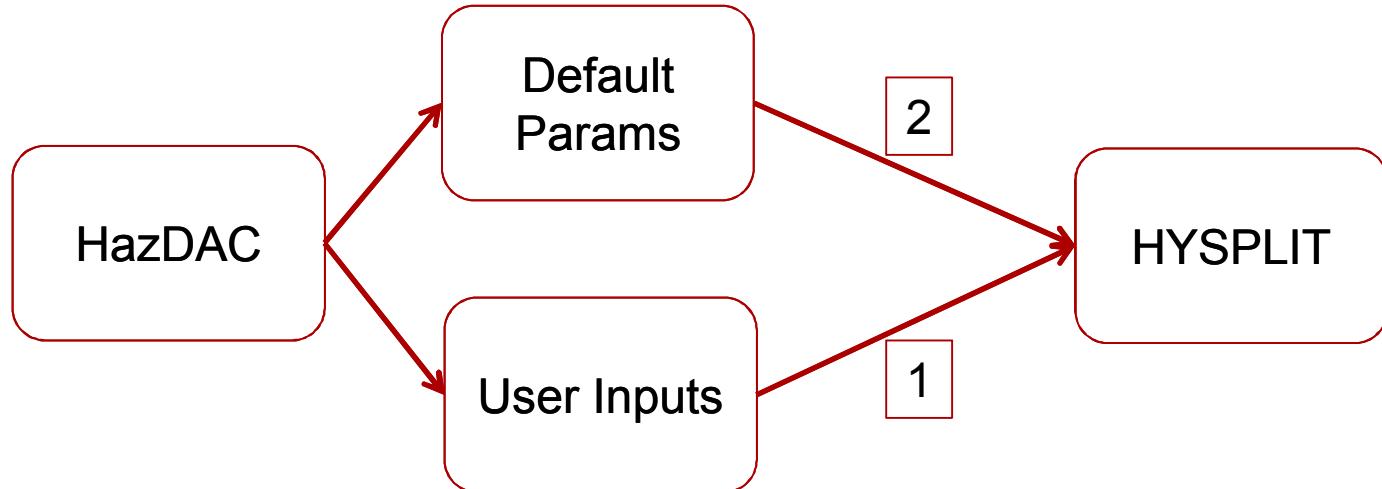
Model Differences Consistent

- HPAC consistently results in higher probabilities of detection



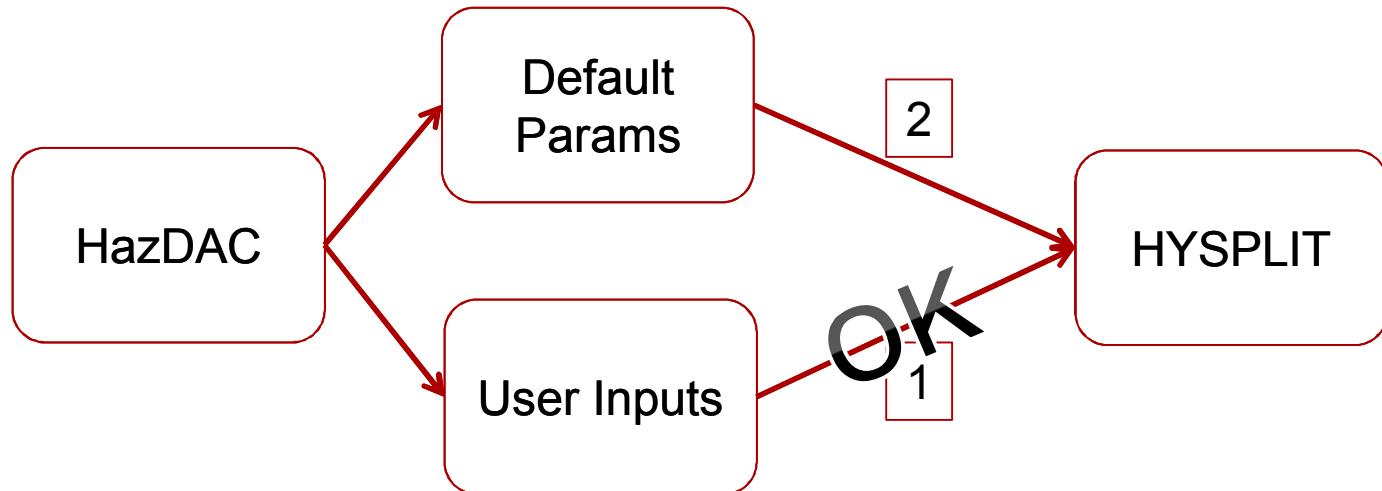
Three Possible Explanations

- 1) HazDAC is not translating user inputs to HYSPLIT correctly
 - Can be easily fixed
- 2) Default parameters in HYSPLIT are different than HPAC
 - Can probably be identified and fixed
- 3) HPAC and HYSPLIT are fundamentally different
 - Experimental validation required



1) Translating User Inputs: OKAY

- Inputs such as mass released, release location, etc.
- Conclusion: User inputs correctly given to HYSPLIT



2) HYSPLIT Default Parameters

- Parameters that can be easily changed:
 - Grid Spacing
 - Time step
 - Release particle number
 - Gaussian distribution
- Other advanced parameters:
 - Deposition constants
 - Split-merge constants
 - Turbulence methods

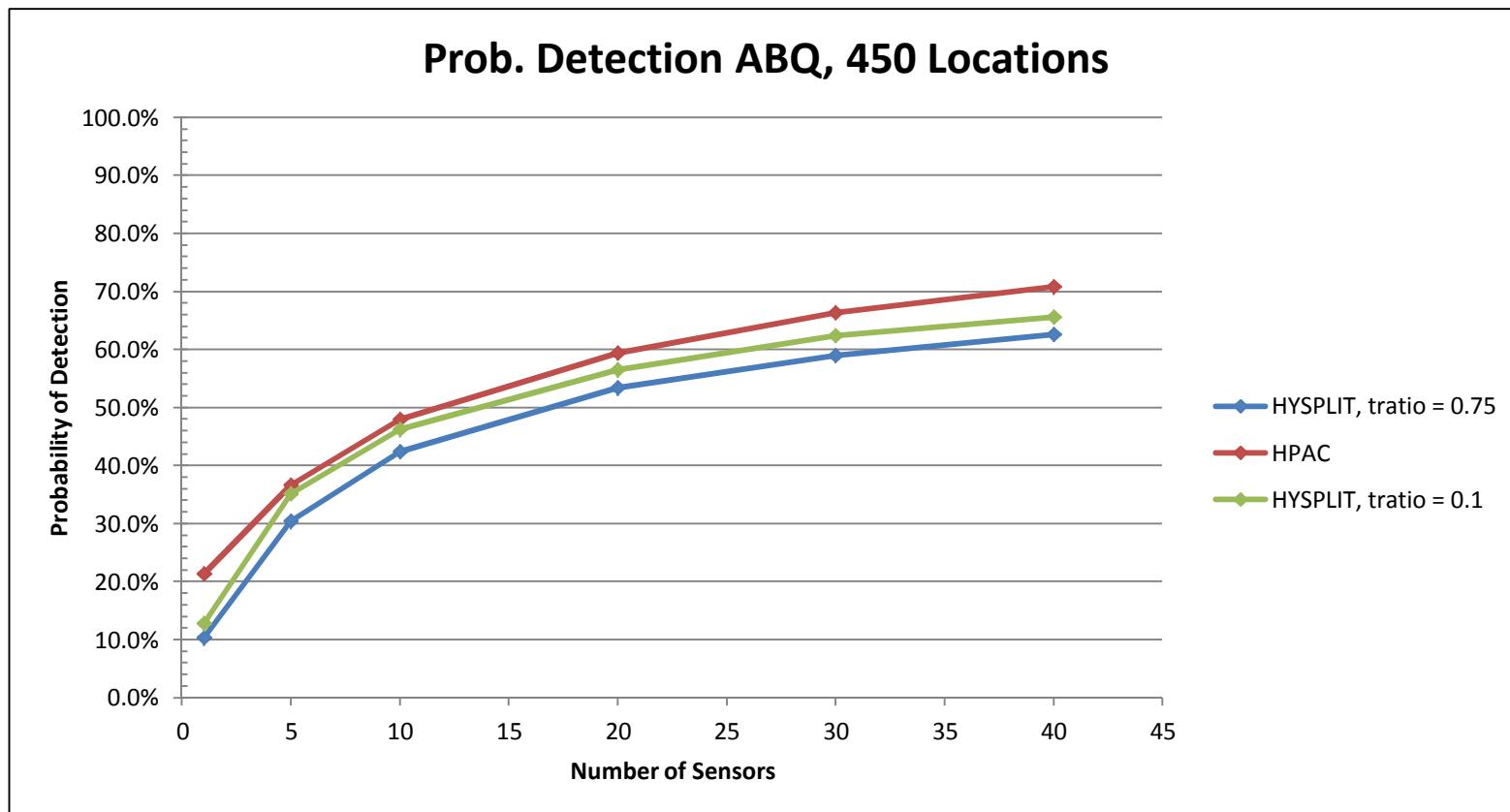
Smaller Grid Reduces Differences

- Decreasing grid size brings HYSPLIT curve close to HPAC



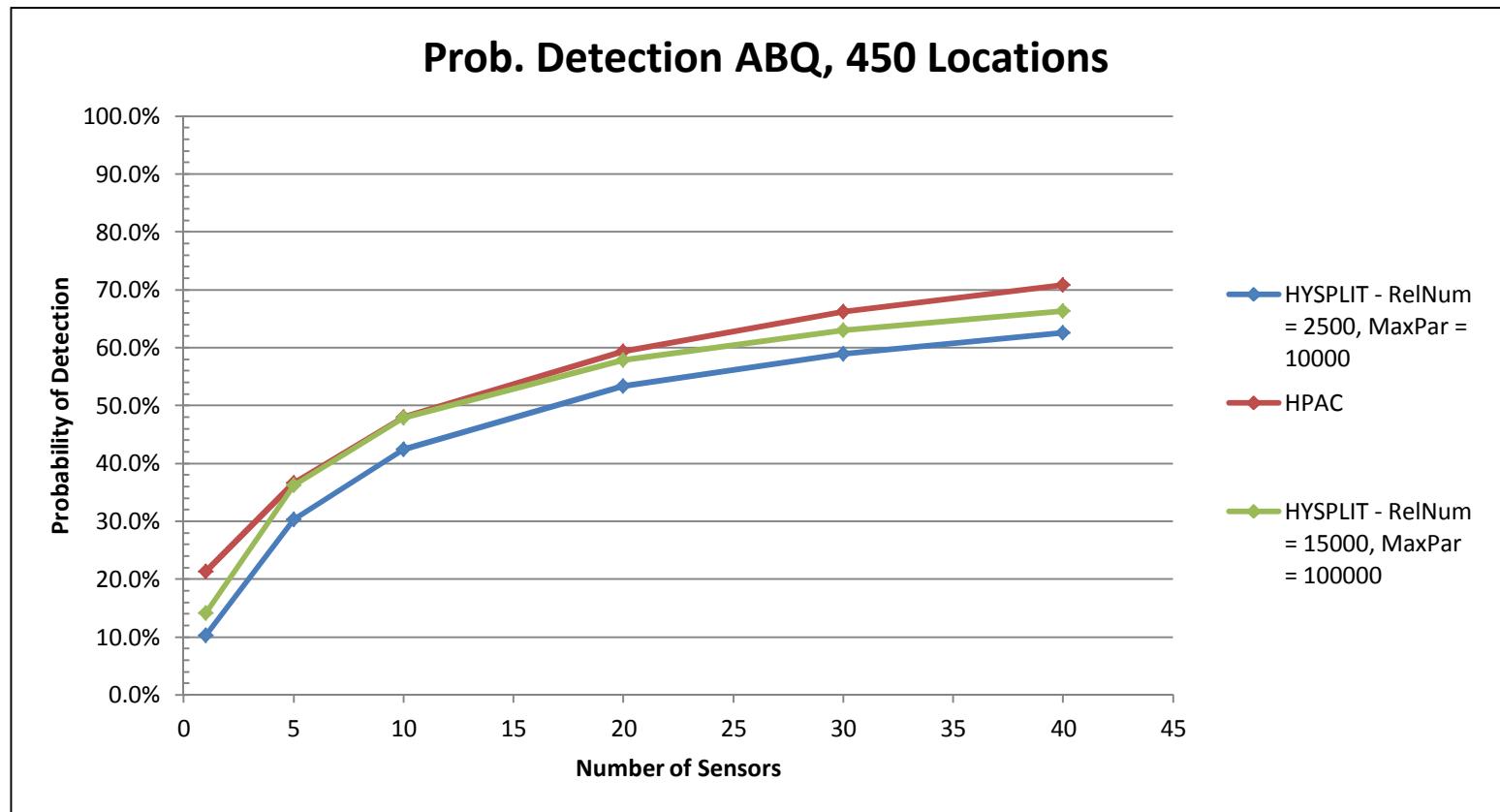
Smaller Δt Reduces Differences

- Decreasing time step brings HYSPLIT curve closer to HPAC



Larger Releases Reduce Differences

- Increasing the number of initial particles released brings curves closer

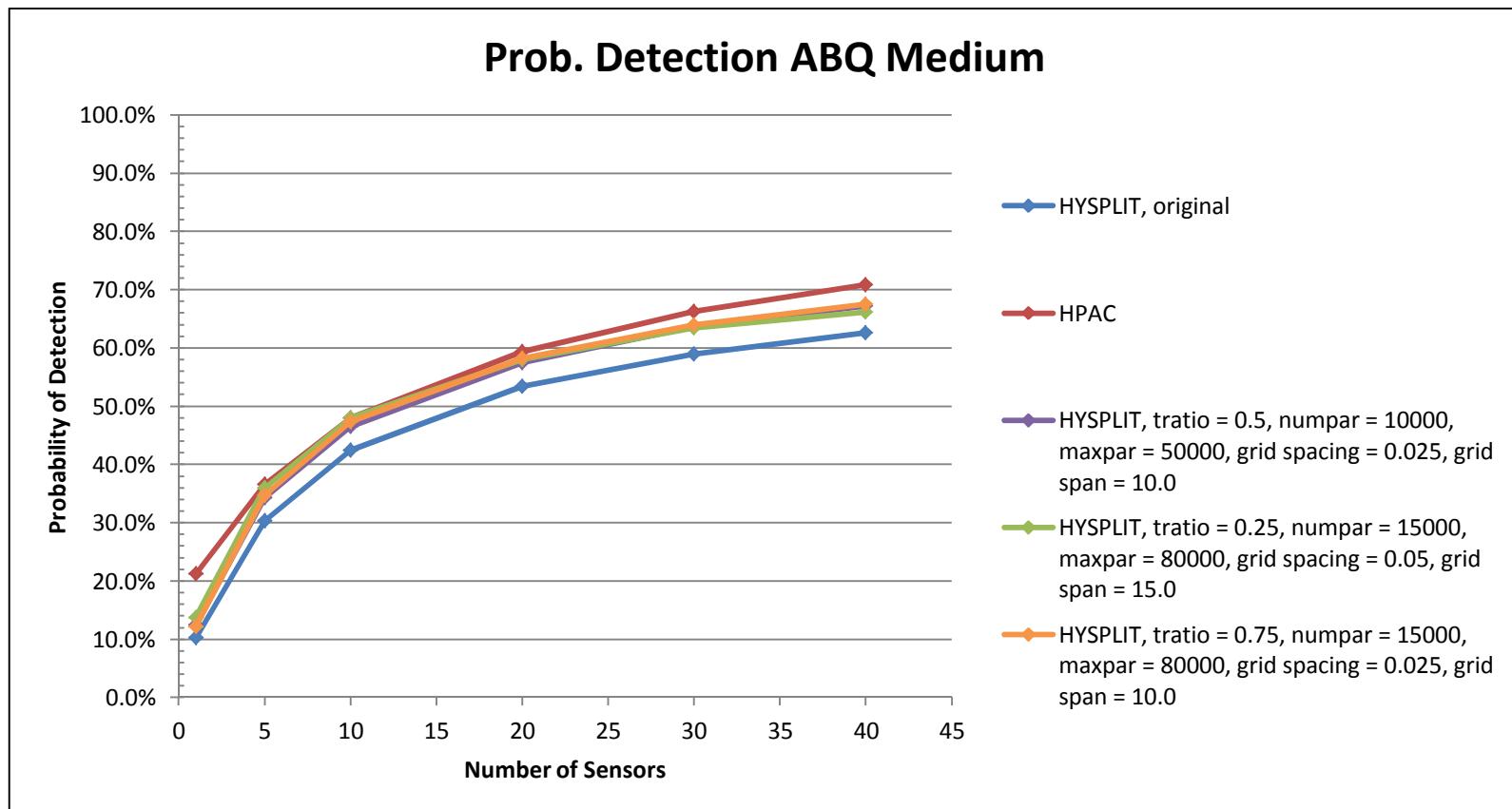


Time/Memory vs. Accuracy

- Important parameters that affect accuracy
 - Grid spacing, time step, and release numbers
- Smaller time step = more accurate, but also longer
- Same with grid size and release numbers
- Where is the happy median?
 - HPAC's exact parameters are unknown

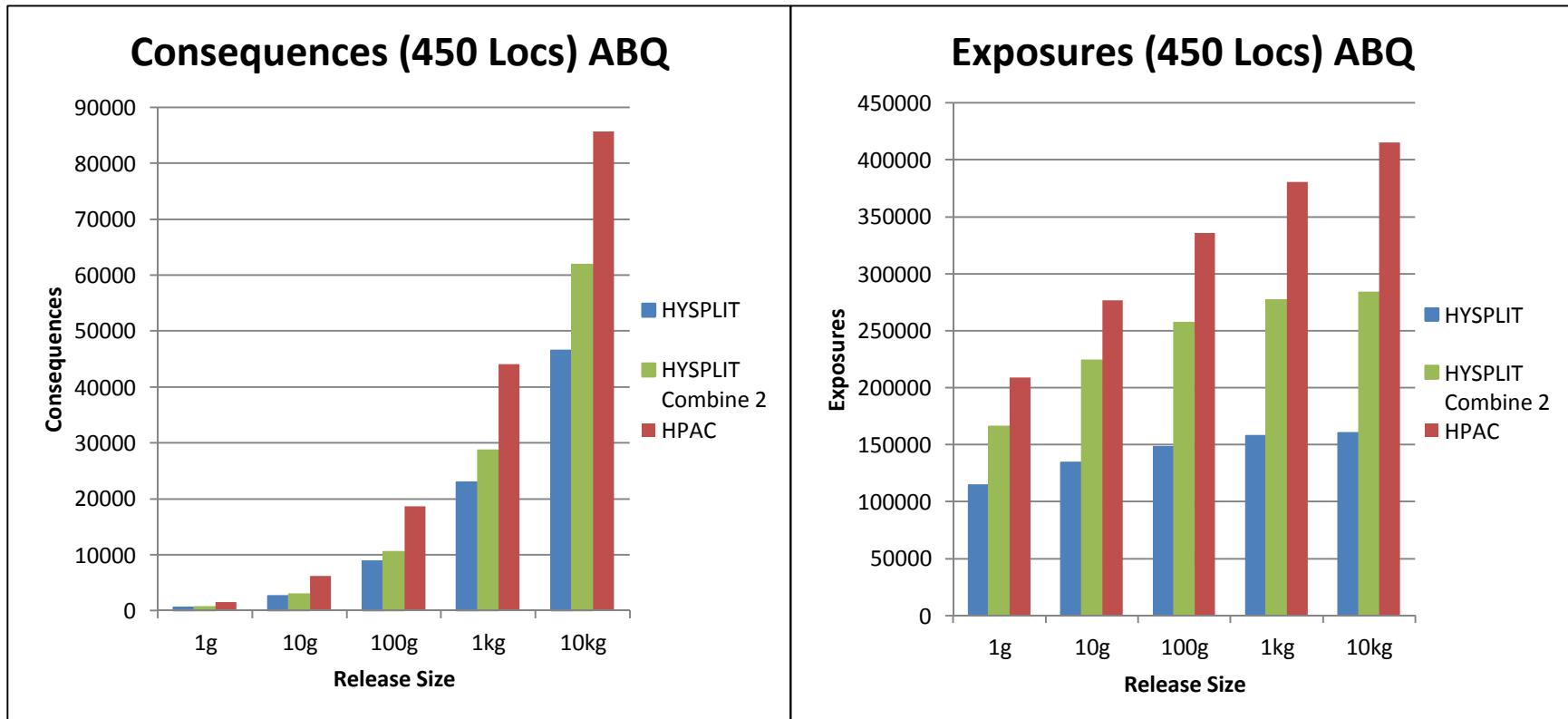
Happy Median: Δt and Release Size

- Minimal compounded effect
 - Release number has greatest impact



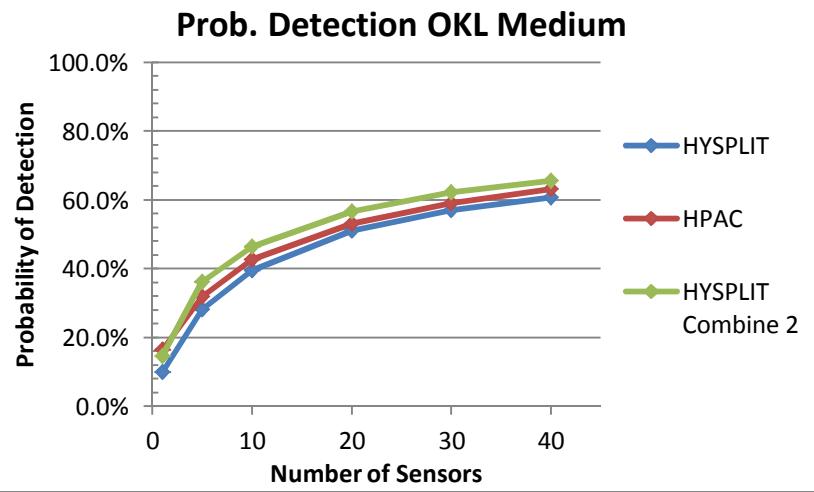
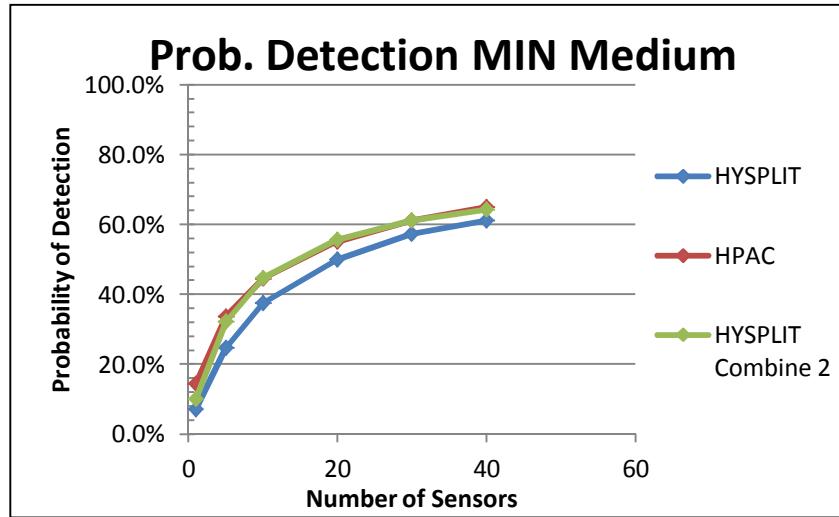
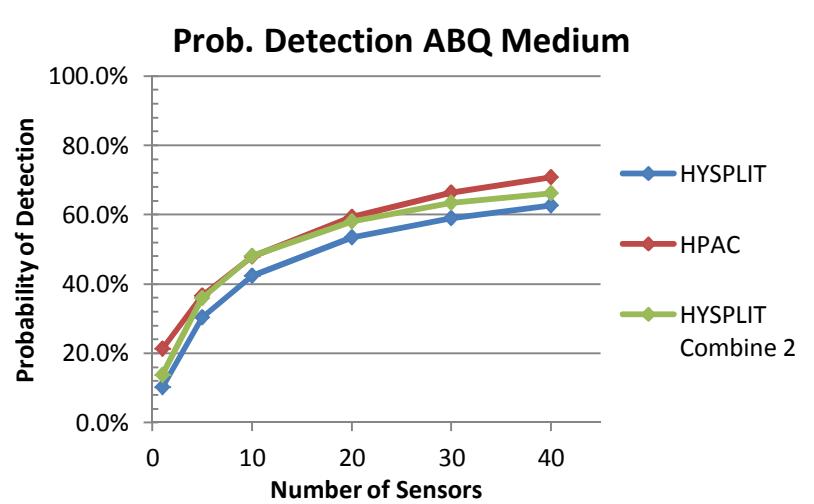
Health Predictions Comparable

- HYSPLIT numbers much closer to HPAC numbers
- HPAC still consistently results in larger numbers



Similar Results in Other Cities

- HYSPLIT curves match HPAC ones very well!
- Successful adjustments



HazDAC Summary

- In general, HYSPLIT gives lower predictions
- Three possible parameters to bring numbers closer
 - Implementation of HYSPLIT in code - OKAY
 - Default parameters for HYSPLIT – Some Problems
 - Fundamentally different models – Still a possibility
- Release numbers are most important factor
- But, HPAC still has generally higher numbers than HYSPLIT
- Likely, the models are fundamentally different and will never yield exactly the same numbers

Future HazDAC Work

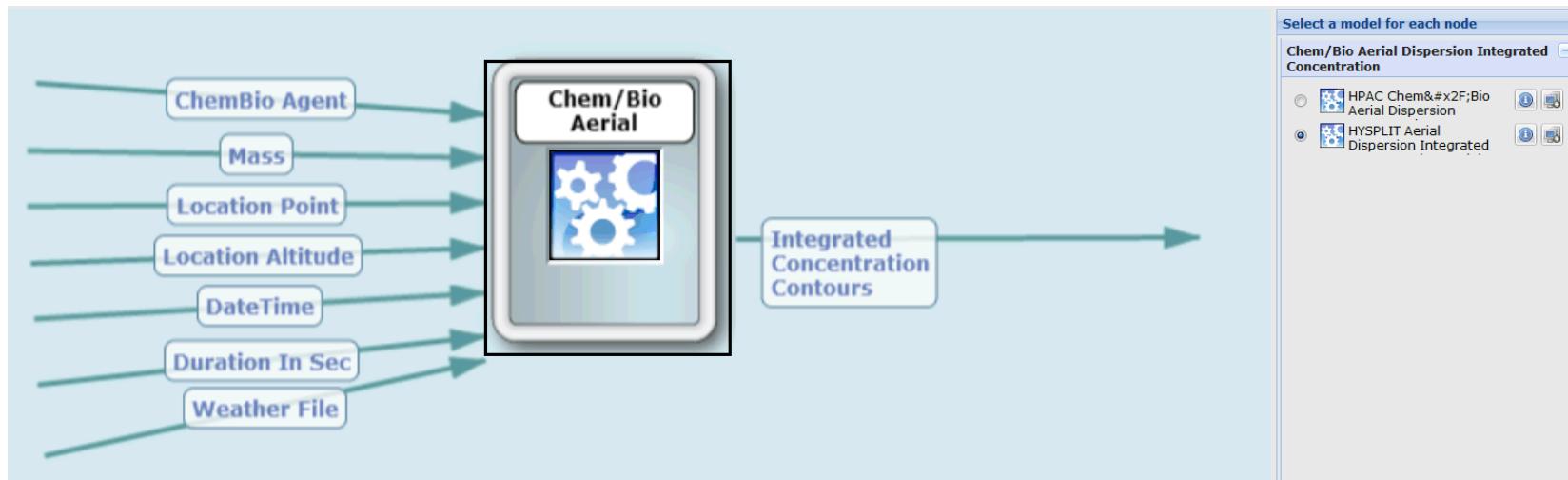
- Find absolute best happy median
 - Requires more runs with variations in parameters
- Contact expert in HPAC (and maybe HYSPLIT)
 - Figure out *exactly* what the differences are
- Run more tests with different agents, cities, etc.
- Validate against real life tracer experiments
 - This could demonstrate whether HPAC or HYSPLIT is “better”

SUMMIT: Background

- SUMMIT links different models together
 - Has models for many purposes
- Goal:
 - To give users the most options
 - To pull together similar software in one place
- SUMMIT will model aerosol dispersion for chem/bio agents
 - Needs to incorporate HYSPLIT

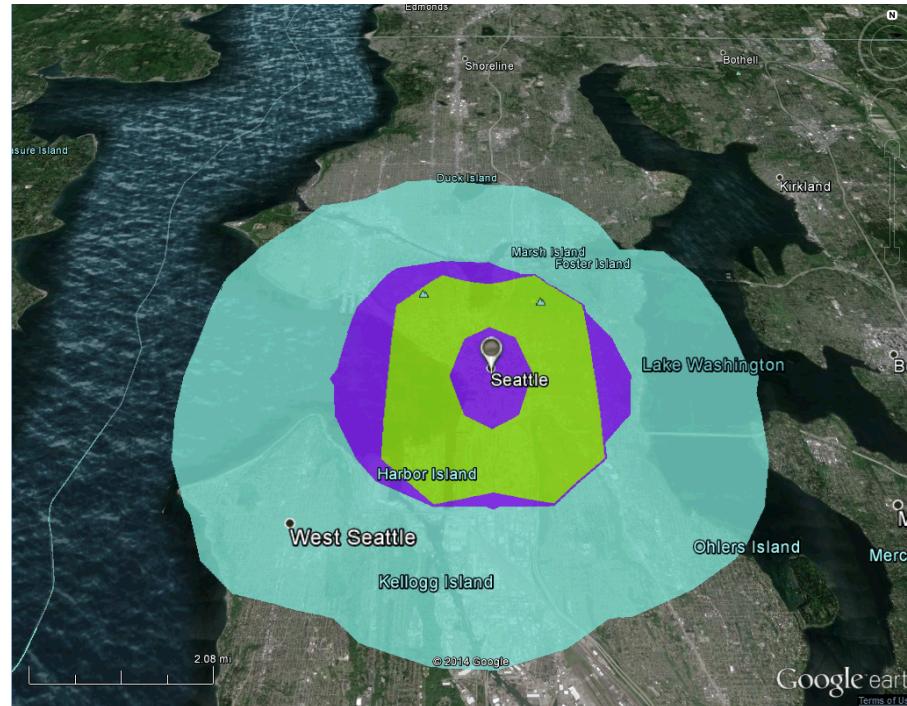
HYSPPLIT in SUMMIT

- Want HYSPPLIT to be correctly implemented
- Want inputs to be the same as for similar models (i.e. HPAC)
- Want outputs to be displayed in user-friendly format



Success!

- Used knowledge from HazDAC discussed above
- Implemented of HYSPLIT for releases
- Wrote implementation code



Future SUMMIT Work

- Document thoroughly for future coders
- Run more verification tests
 - SUMMIT is different use case than HazDAC
- Implement a time-lapse output to see plume moving
- Add other more specific inputs for more advanced users

Conclusions

- Bioterrorism is bad.

Conclusions

- Bioterrorism is bad.
- But Sandia (and California) is fun!

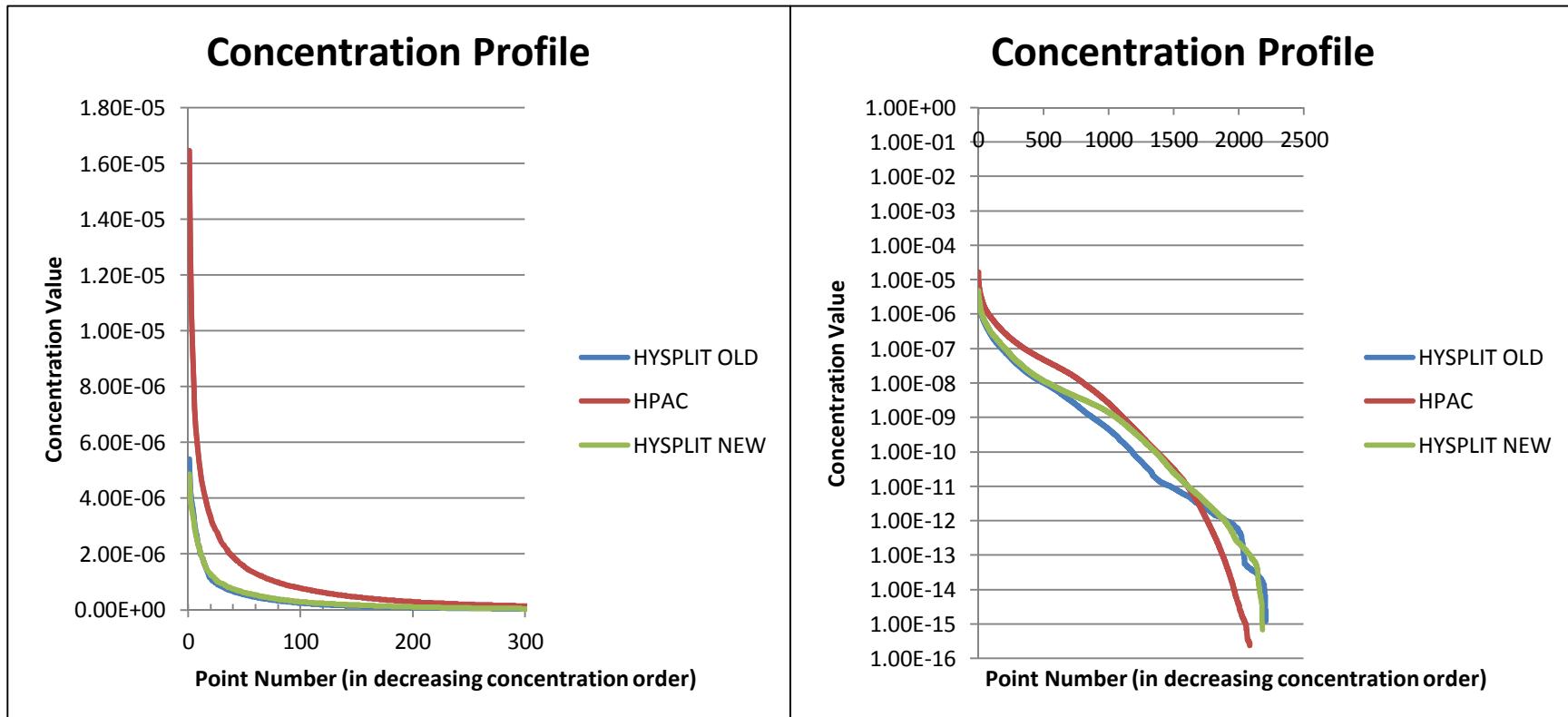
PICTURE OF INTERNS HAVING FUN HERE

Thanks!

- Thanks to my mentors Dr. Peterson and Dr. Teclemariam
- Thanks to all the other mentor-like people
- Thanks to the interns for being generally awesome
- Thanks to Sandia for giving me this opportunity

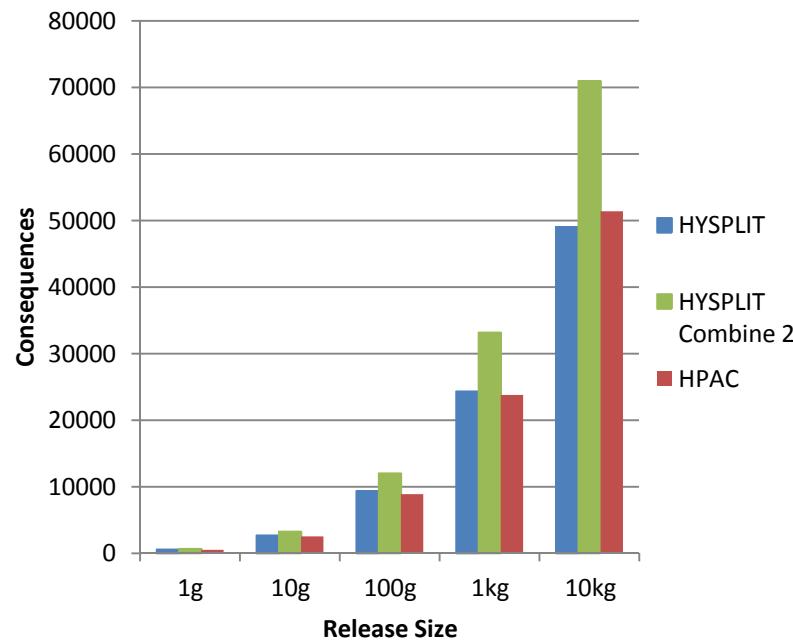
Improved Concentration Profiles

- No obvious improvement...
- But could be grid spacing? Or just ABQ?

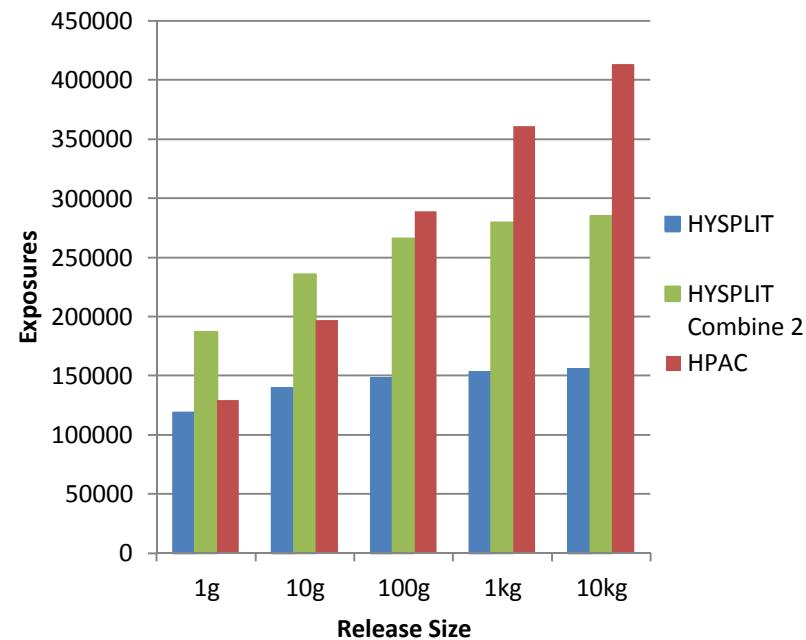


Health Effects: OKL

Consequences (440 Locs) OKL



Exposures (440 Locs) OKL



Health Effects: MIN

