

Report 12/11/2020

P. Hacker

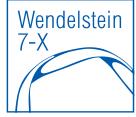


Previously: 2D Inversion Results



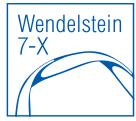
- considering a given emissivity profile shape, there exists an ideal set of K factors constraining the ratio of poloidal and radial gradients according to the χ^2 optimisation
- not necessarily the same as defined by the lowest 2D deviation and Pearson coefficient, whose optimisation also yields a set of K factors
- radial positioning of profiles is subject to the selection of K factors over the radius (and angular space)

Previously: 2D Inversion Results



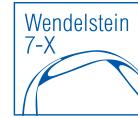
- poloidally distributed profiles are, so far the data set suggests, the hardest to reconstruct properly (especially on the inboard side)
- asymmetric characteristics can be property of the underlying emissivity profile to the tomography, however some are introduced due to the line of sight geometry
- combinations of profile shapes, especially poloidally symmetric and any else become hard to distinguish from one another (see ring and island mimic)

Previously: 2D Inversion Results



- bright spots on-/off-axis and, e.g. bright rings can be separated from each other depending on the distance
- individual bright sheaths can no longer be reconstructed radially when closer than 0.2m
- reconstructions of forward calculated STRAHL simulations are largely accurate in position and value
- reference XP 20181010.032 difficult to reconstruct, subject to difference in forward and backward model?

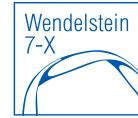
QSB LabVIEW Routine Review (see CHANGELOG.md)



PROS

- added EPOCH timer to UI, timing mode that calculates EPOCH time at start
- added T1-T4 timing input to UI, added T0-T1 timer to UI (should at least be 30s)
- fixed wrong time base in calibration procedure
- DAC range setting file load or input via the UI
- **selected channel feedback output**, though with delay >10ms (single/multi channel)
- performance for 1-10 selected channels in feedback is fine
- overhaul of calibration parameter routines since detector response is more complex than a single peak and exponential decay
- full P_rad routine added for **after** the data acquisition to access power loss quickly (same as IDL/python)
- filter register settings fixed to match sample timings in milliseconds
- improved/implemented local-storage-mode-only, so no crash at upload
- upload of error corrected voltage input to archive as well (same as IDL/Python)

QSB LabVIEW Routine Review (see CHANGELOG.md)



CONS

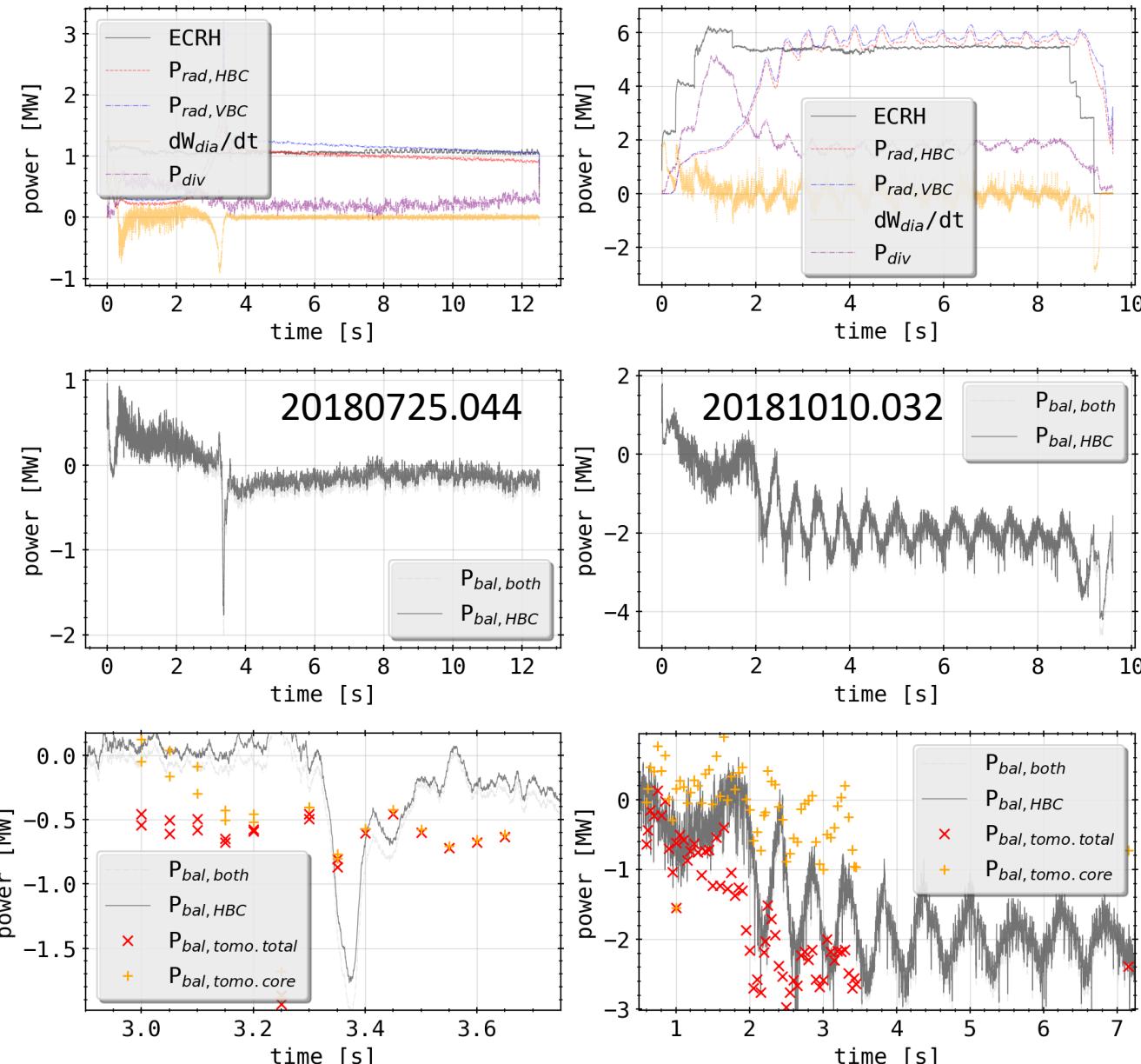
- DAQ routine now more prone to crashing, hiccups and **slow processing causing sample timing errors**
- ADC bit settings not fully understood, potentially not intended setting here (wrong documentation?)
- crashing with bad device status at memory overflow or uncompleted DAQ/upload, inconsistent to recover
- hardware needs warm-up time of ~30min (before?)
- some small fixes needed to improve quality-of-life experience during experimental campaign (dials not working as intended, ...)

Tomogram Statistics: Experiment Power Balance

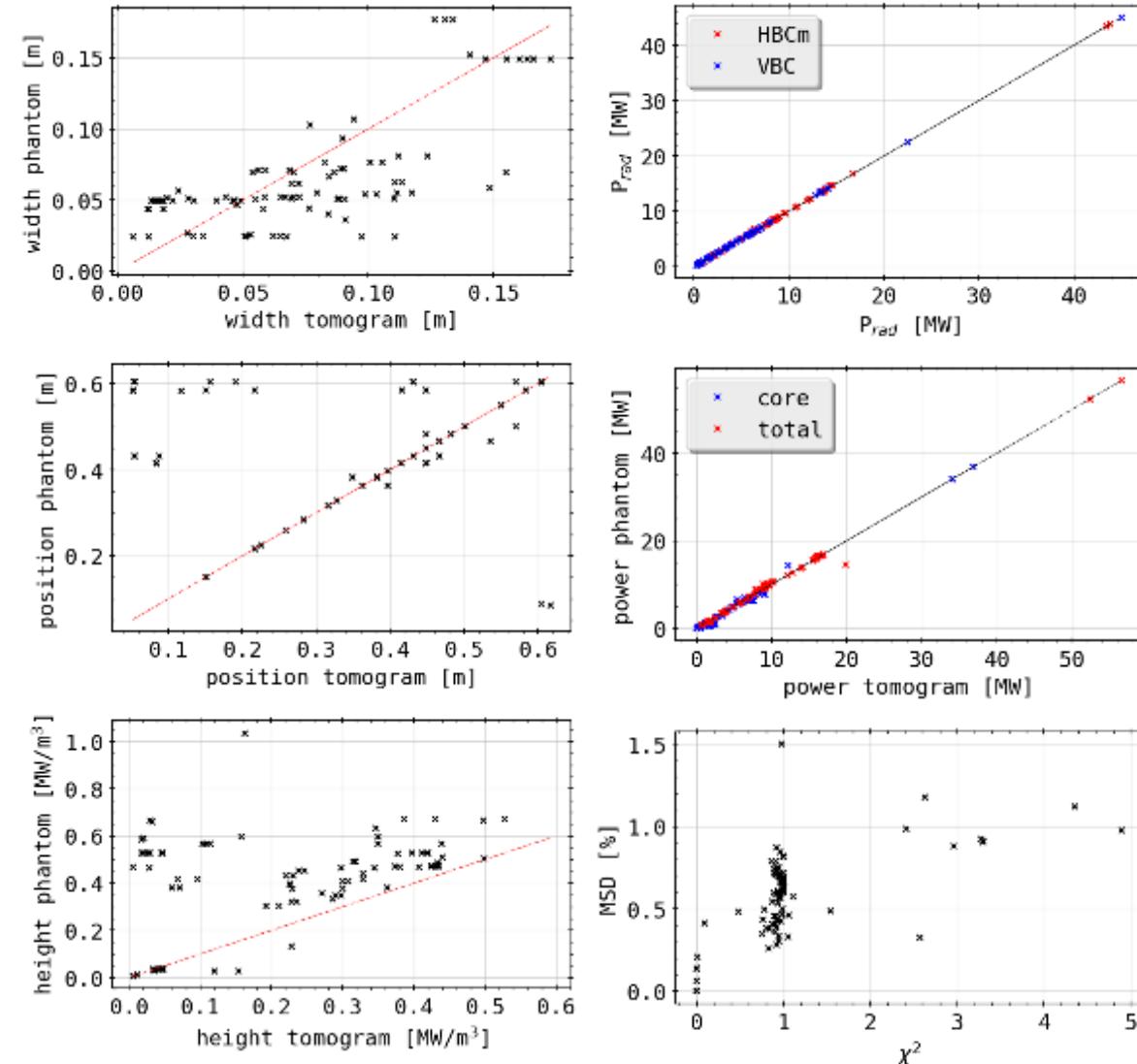
- compiled experiment data and parameters to calculate

$$P_{bal}(t) = P_{ECRH}(t) - P_{rad,HBC/VBC}(t) - P_{div}(t) - \frac{dW_{dia}}{dt}(t)$$

- also exchanged radiated power proxy from bolometer camera by 2D integrated power of tomogram (both core and total)

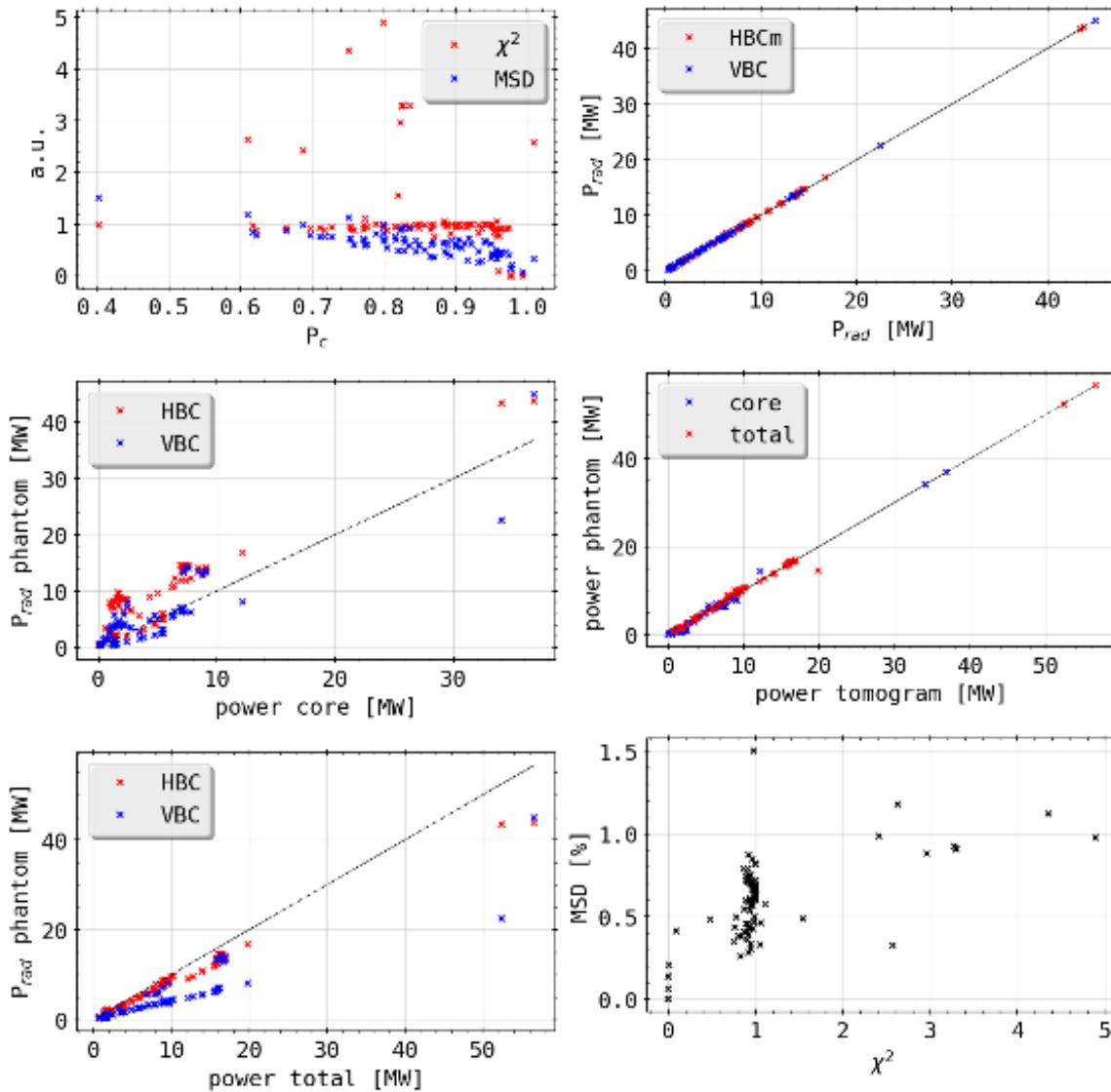


Tomogram Statistics: Phantom Reconstructions



- radial profile positioning good, though width and height have small to large deviations
- peak height comparison exhibits offset on phantom side: widened radial shape and longer decays (width)
- P_{rad} and total/core power values match very well for phantom reconstructions
- a slightly larger range of MSD values yields a ‘good’ χ^2 of ~ 1.0
- there seems to be an offset of MSD about 0.25% (minimum error between source and target to be expected?)

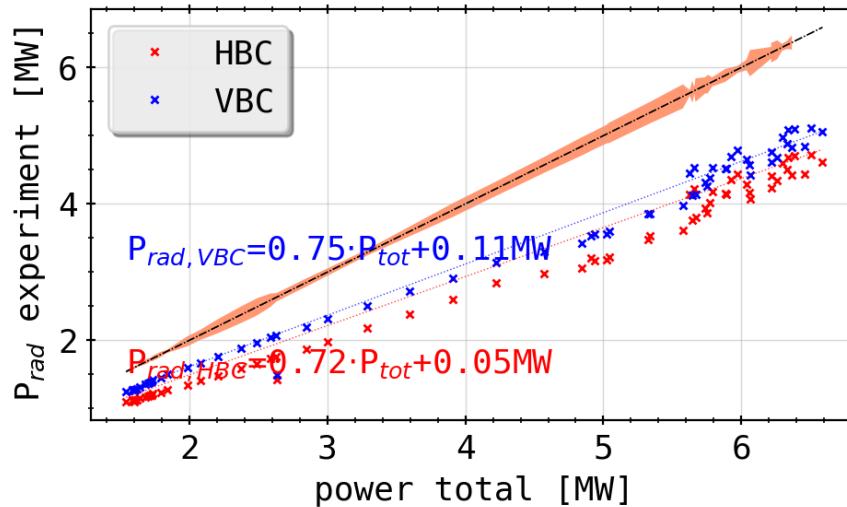
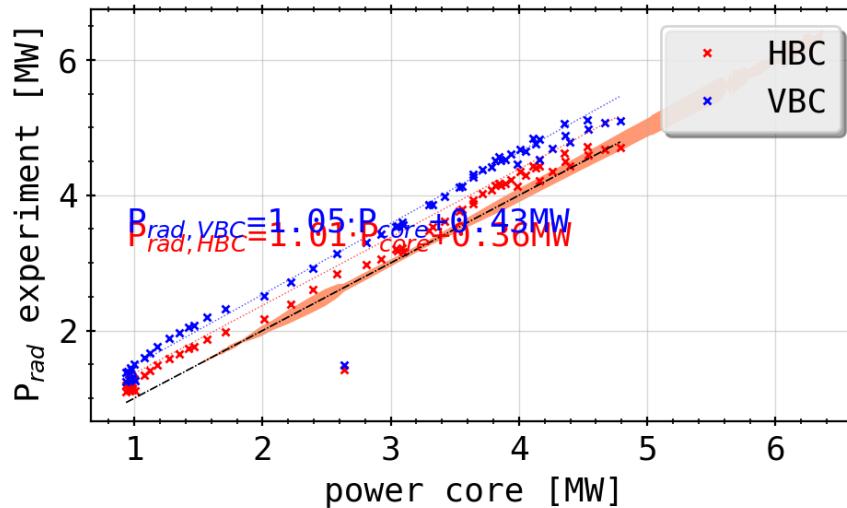
Tomogram Statistics: Phantom Reconstructions



- MSD approaches 0.0 if pearson coefficient reaches 1.0, as expected, while χ^2 is around 1.0
- P_{rad} over core/total 2D integrated power yield constant offsets and slopes
- core power matches P_{rad} vor VBC(?) though is lower for HBCm as expected
- total power is larger for both P_{rad} estimates, though with different proportionalities

Tomogram Statistics: Experiment Reconstructions Errorbar

20181010.032



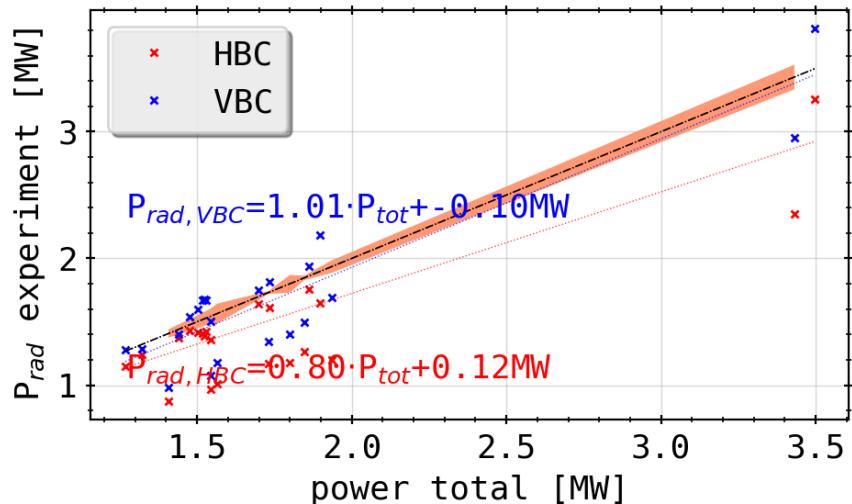
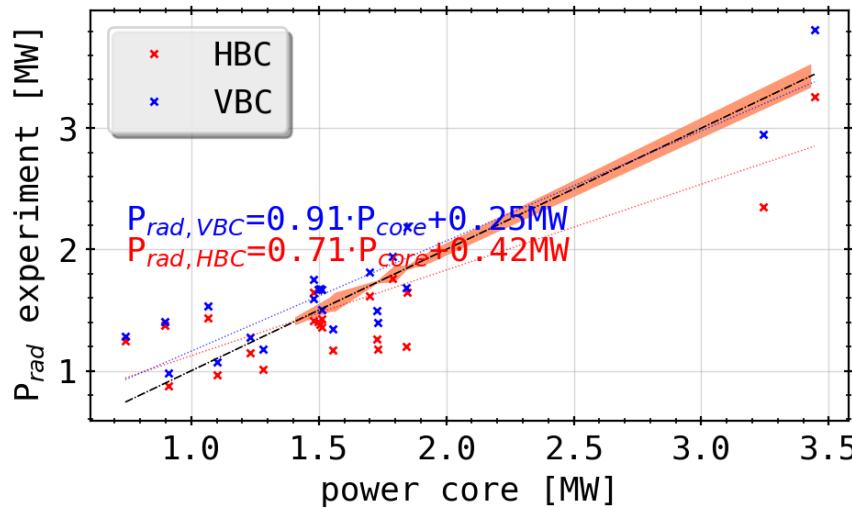
- extra slope in P_{rad} vs. total power by cutting negative results from reconstruction when limiting to only physically meaningful numbers ($P/m^3 > 0$)?
- increase of their impact is expected with overall higher radiation level to counteract errors in chordal profile

CONCLUSION

- impact of mathematical error is negligible across the power spectrum, but why?
- fit of integrated power values vs camera proxy yields interesting results (~0.4MW offset to core power, ~0.75 ratio to total 2D power)

Tomogram Statistics: Experiment Reconstructions Errorbar

20180725.044

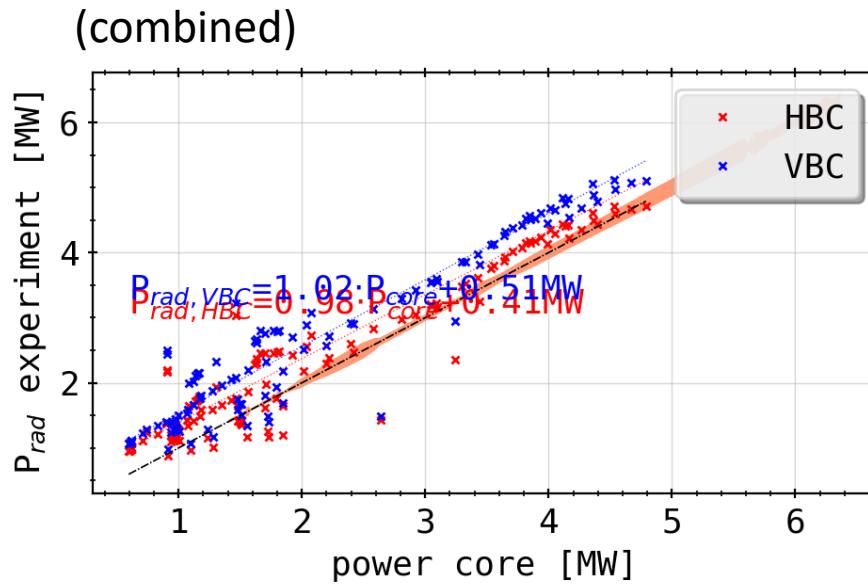


- extra slope in P_{rad} vs. total power by cutting negative results from reconstruction when limiting to only physically meaningful numbers ($P/m^3 > 0$)?
- increase of their impact is expected with overall higher radiation level to counteract errors in chordal profile

CONCLUSION

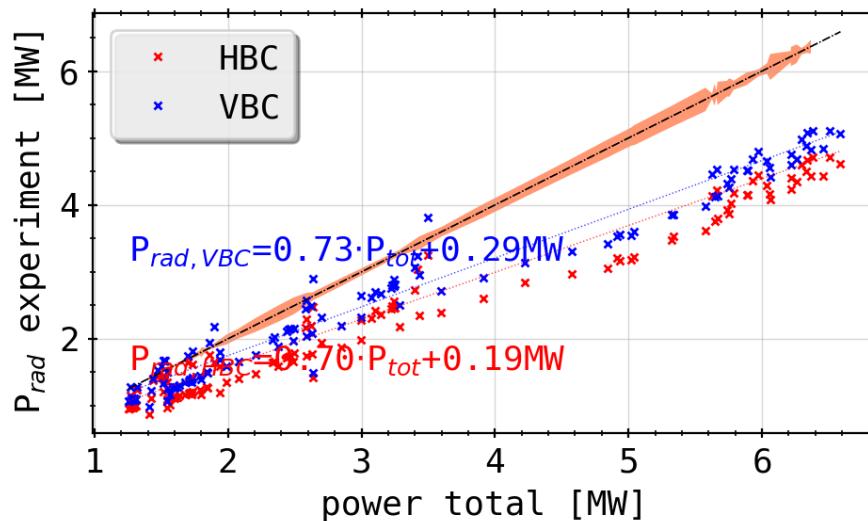
- impact of mathematical error is negligible across the power spectrum, but why?
- fit of integrated power values vs camera proxy yields interesting results ($\sim 0.4 \text{ MW}$ offset to core power, ~ 0.7 ratio to total 2D power)
- changes for different discharge types?

Tomogram Statistics: Experiment Reconstructions Errorbar



- extra slope in P_{rad} vs. total power by cutting negative results from reconstruction when limiting to only physically meaningful numbers ($P/m^3 > 0$)?
- increase of their impact is expected with overall higher radiation level to counteract errors in chordal profile

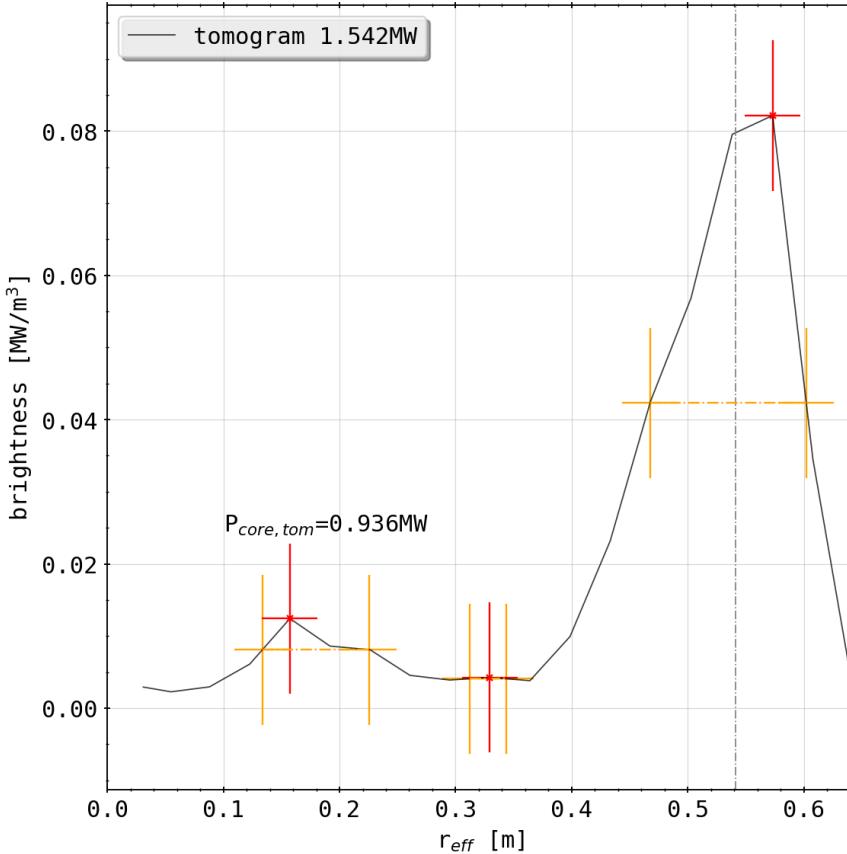
CONCLUSION



- impact of mathematical error is negligible across the power spectrum, but why?
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Tomogram Statistics: Experiment Reconstructions Errorbar

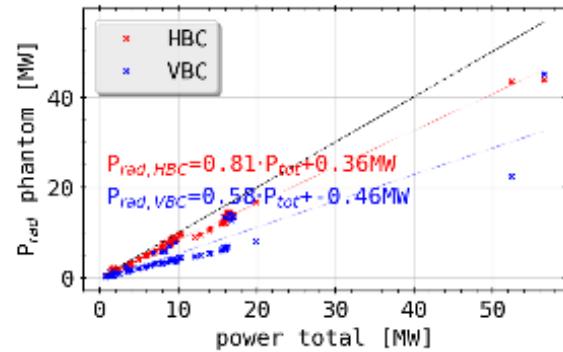
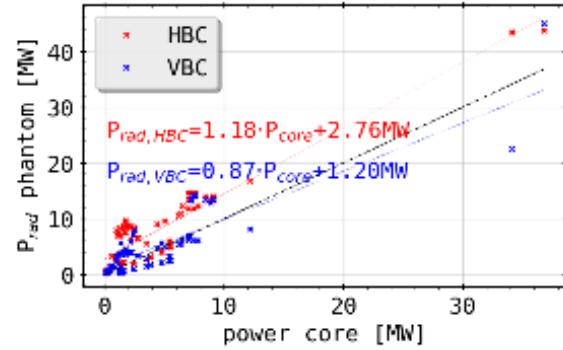
20180725.044@3.45s



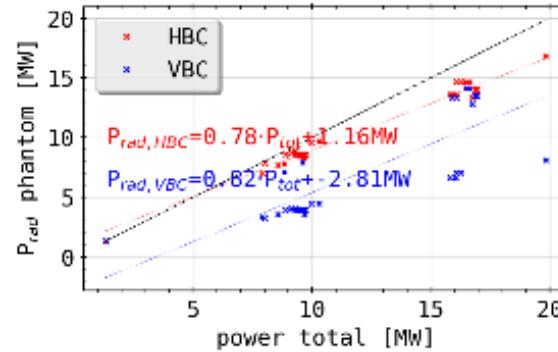
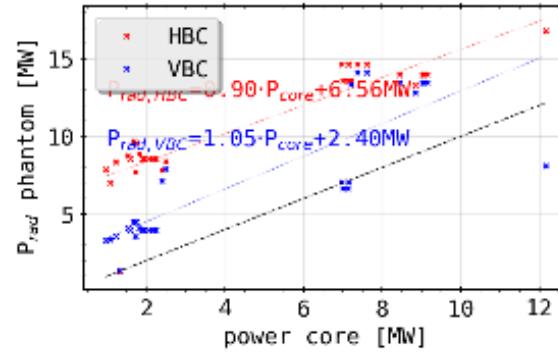
- errorbar looks large here, because the total error for the tomogram (minimum value < 0.0 can be single cell or whole 'ring') has to assumed possible at every integration point

- e.g.: at powers (all $X^2 \sim 1.0$)
total: 5.78 MW , neg: 66.63 kW
total: 6.33 MW, neg: 61.68 kW
total: 2.37 MW, neg: 121.51 kW

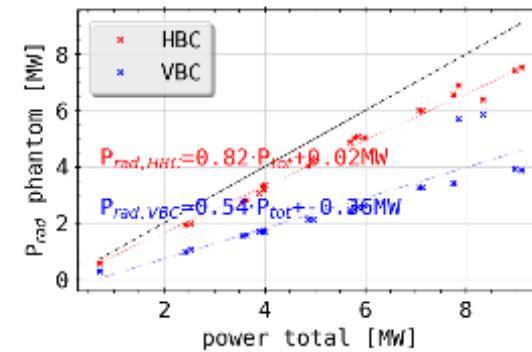
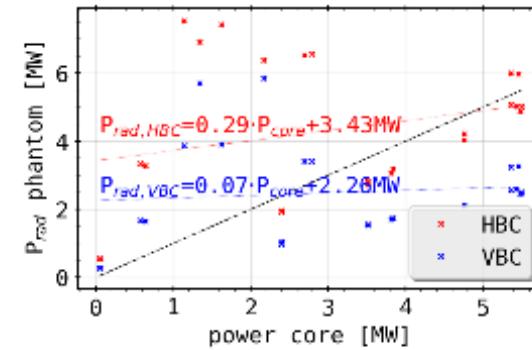
Tomogram Statistics: Phantom Reconstructions



(all phantoms)

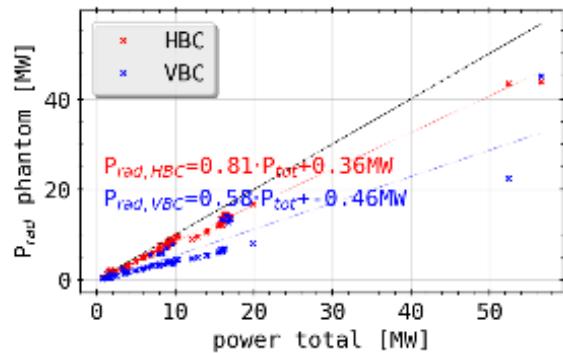
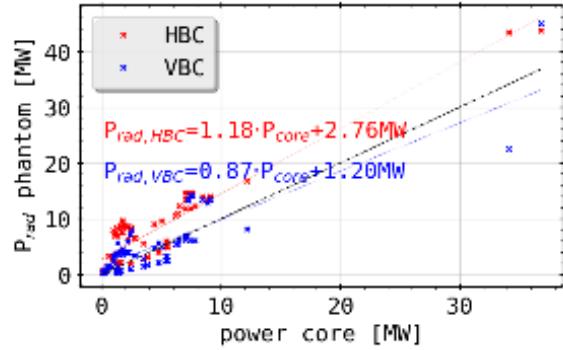


(island mimics)

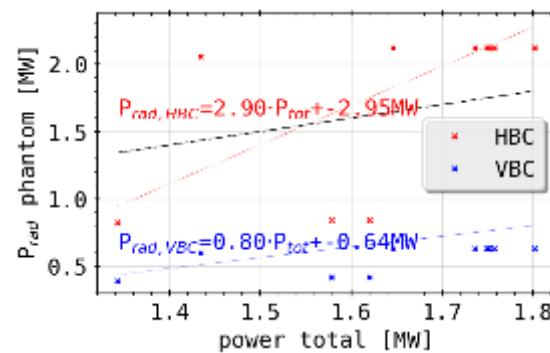
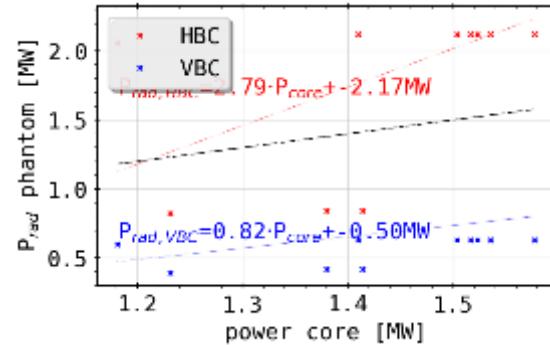


(radiation ring)

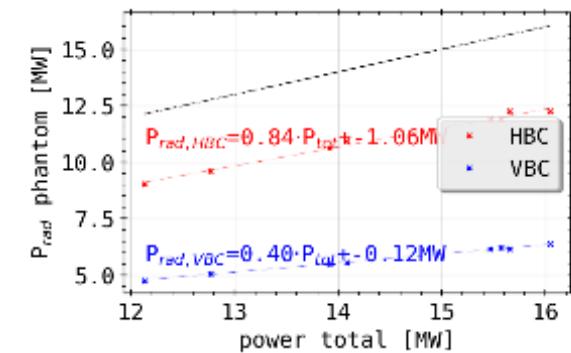
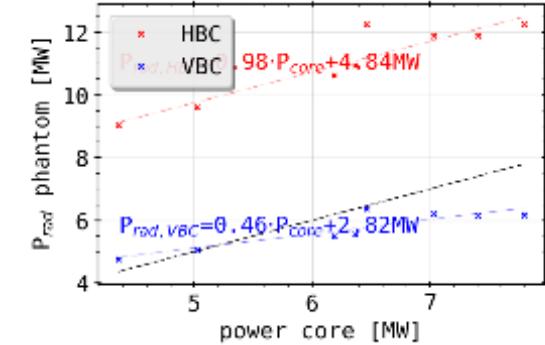
Tomogram Statistics: Phantom Reconstructions



(all phantoms)

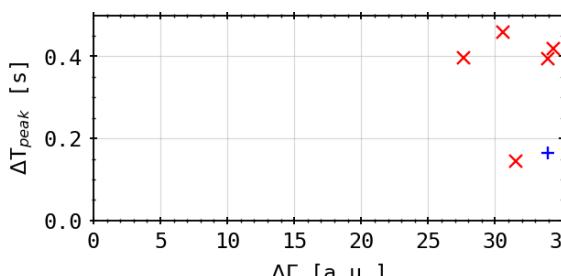
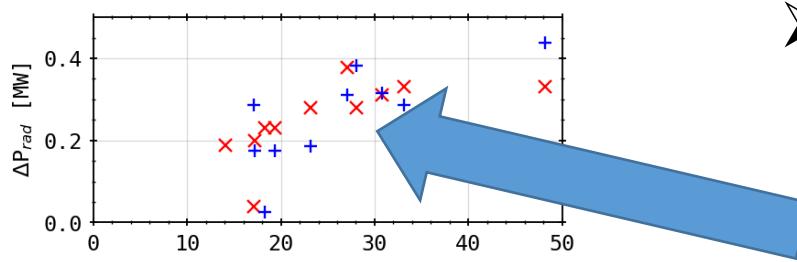
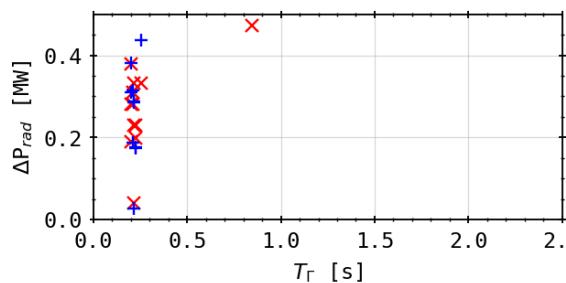
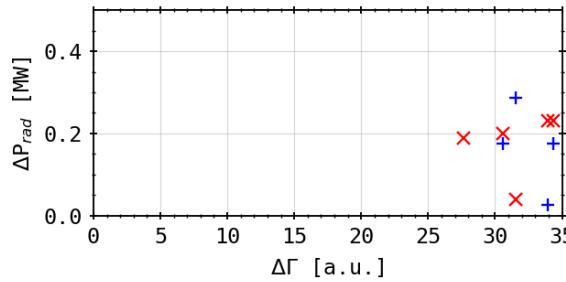


(radiation spot)



(double ring)

Peak Database: Plasma Parameters



➤ results only shown for the feedback controlled discharge
XPID 20181010.032

➤ used *feed-forward* and density control on the main gas valves and the thermal gas valve for radiation loss feedback (H2)

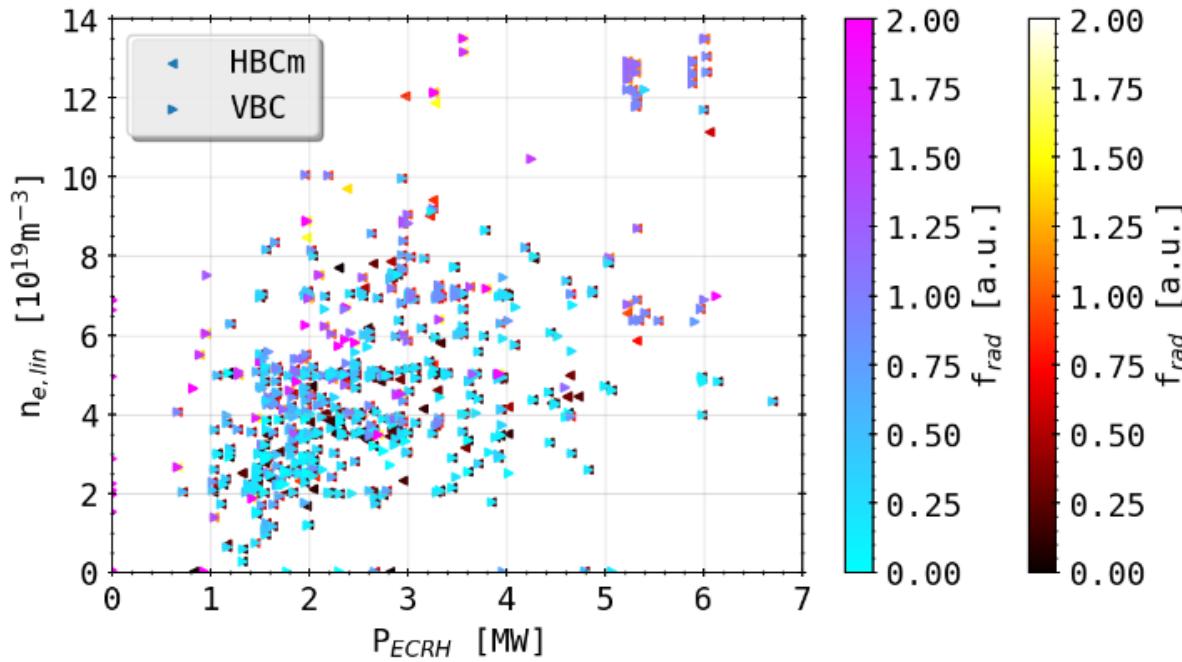
➤ **'fix'** temporal reaction delay of 200ms

➤ linear correlation between puff and radiation reaction:

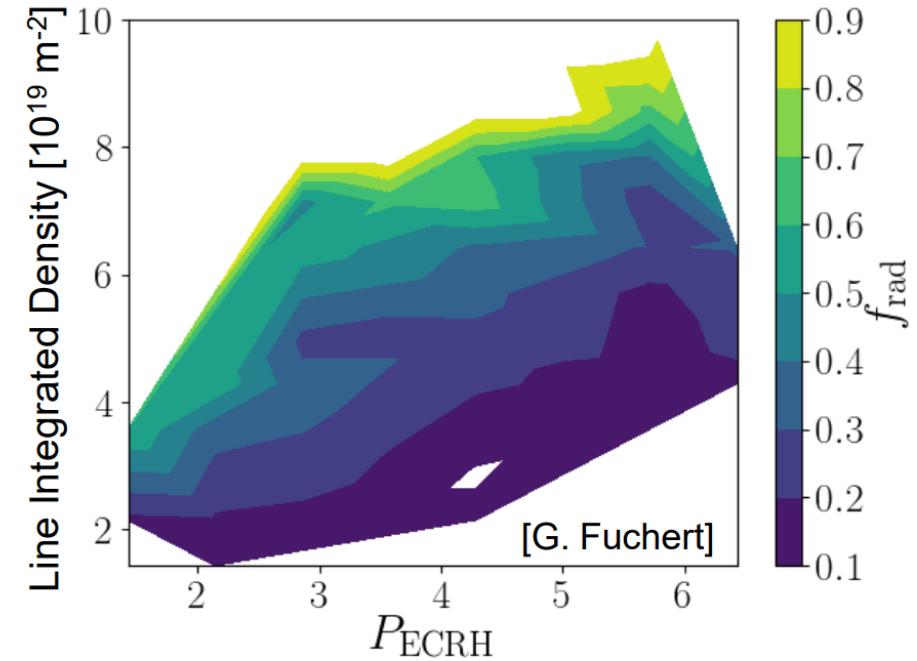
$$\Delta P_{HBCm} [MW] = (0.01187 \pm 0.0063) [MW] \cdot \int \Gamma_{QSQ}$$

$$\Delta P_{VBC} [MW] = (0.00978 \pm 0.0009) [MW] \cdot \int \Gamma_{QSQ}$$

Peak Database: Plasma Parameters



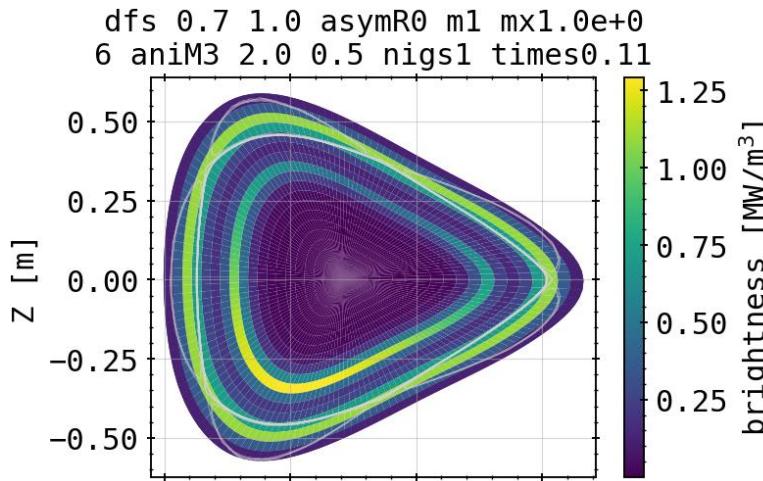
ordering?
→



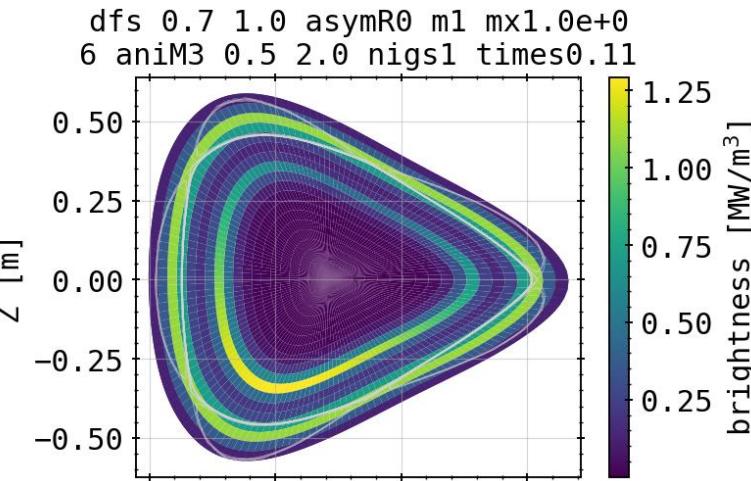
- expecting underlying order of plasma parameters (density, power and exhaust, i.e. energy confinement time)
- further refinement needed: W_dia filtering and confinement time ordering? how when looking at transient phases of gas puff feedback

Asymmetry Propagation: Phantoms

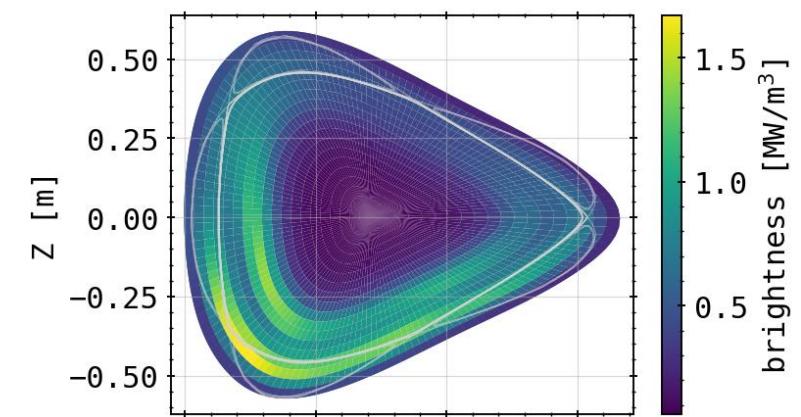
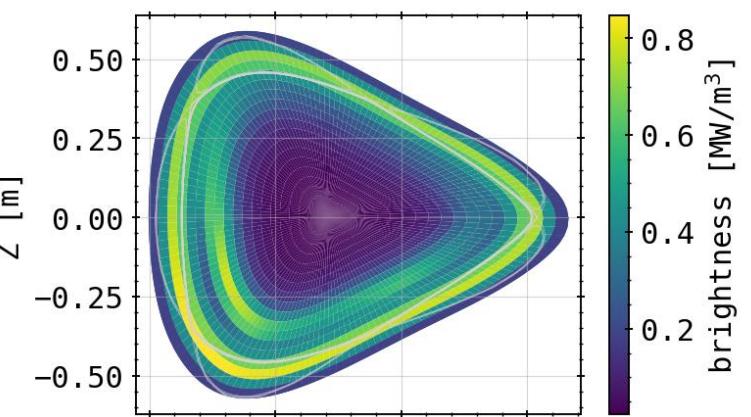
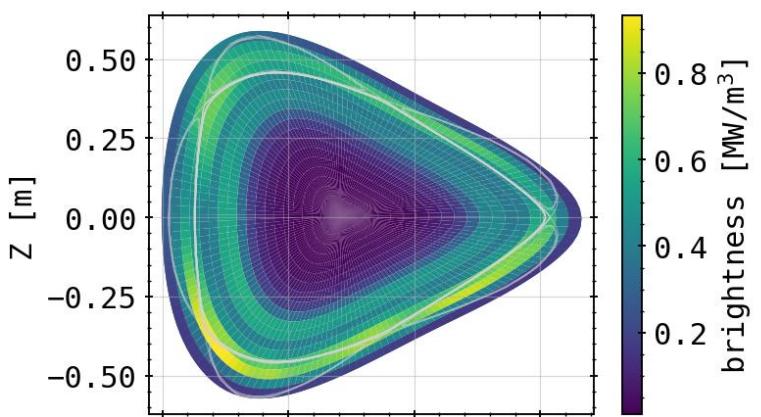
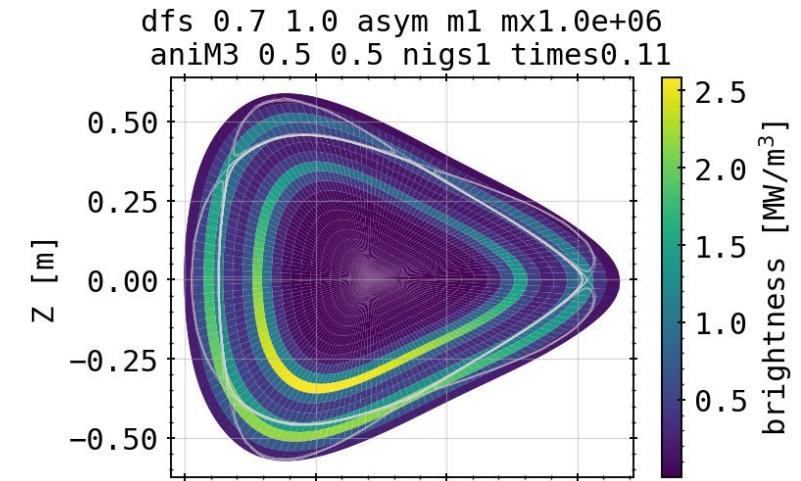
inside asymmetry,
assuming symmetry



inside asymmetry,
assuming asymmetry

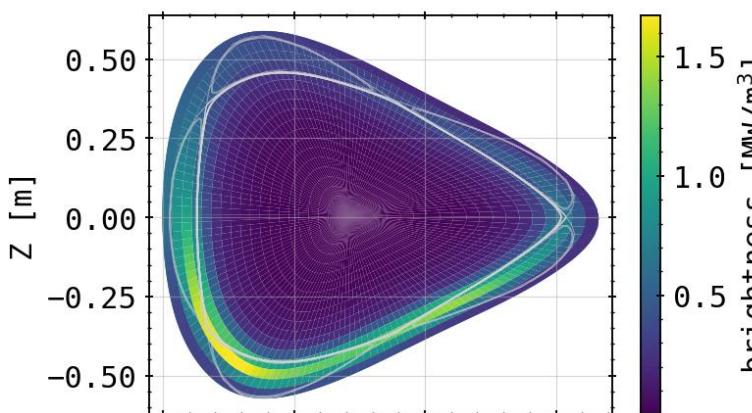
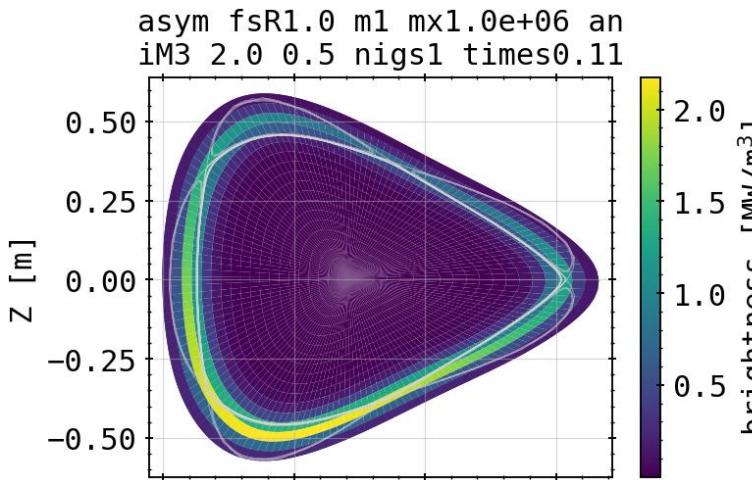


outside & inside asymmetry,
assuming asymmetry

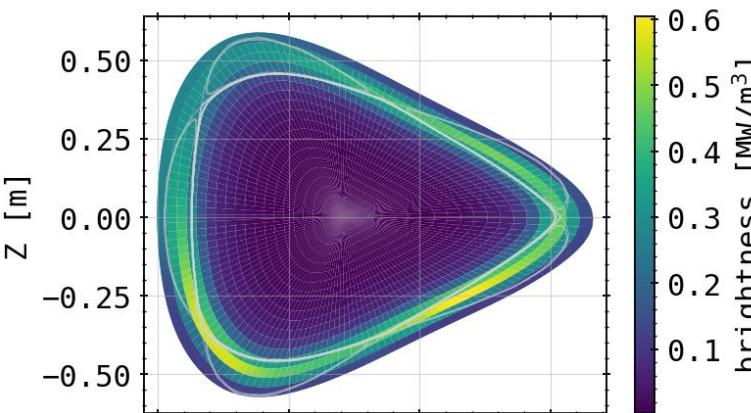
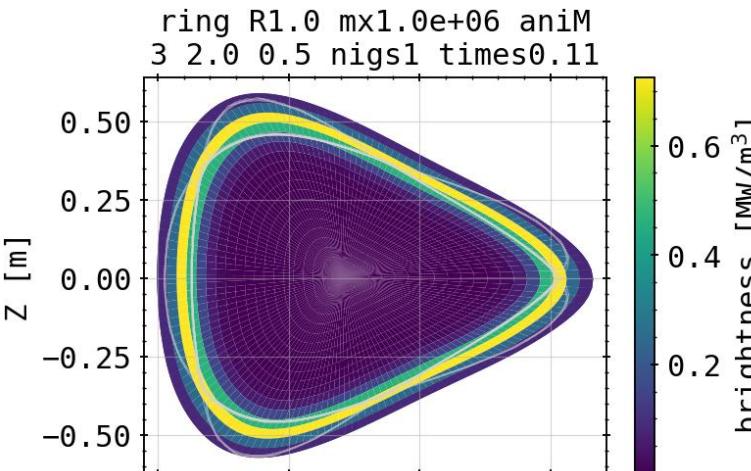


Asymmetry Propagation: Phantoms

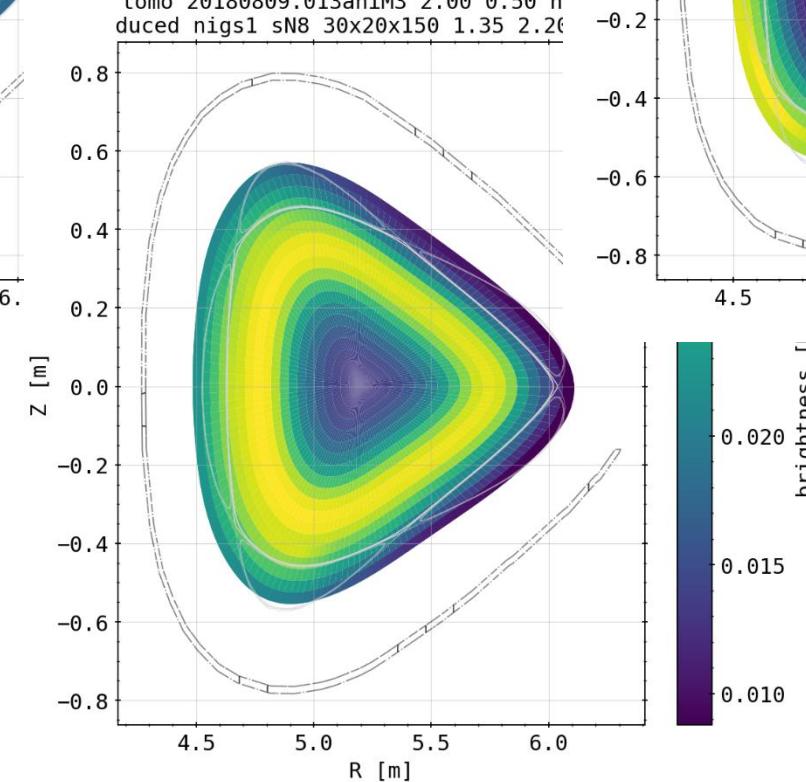
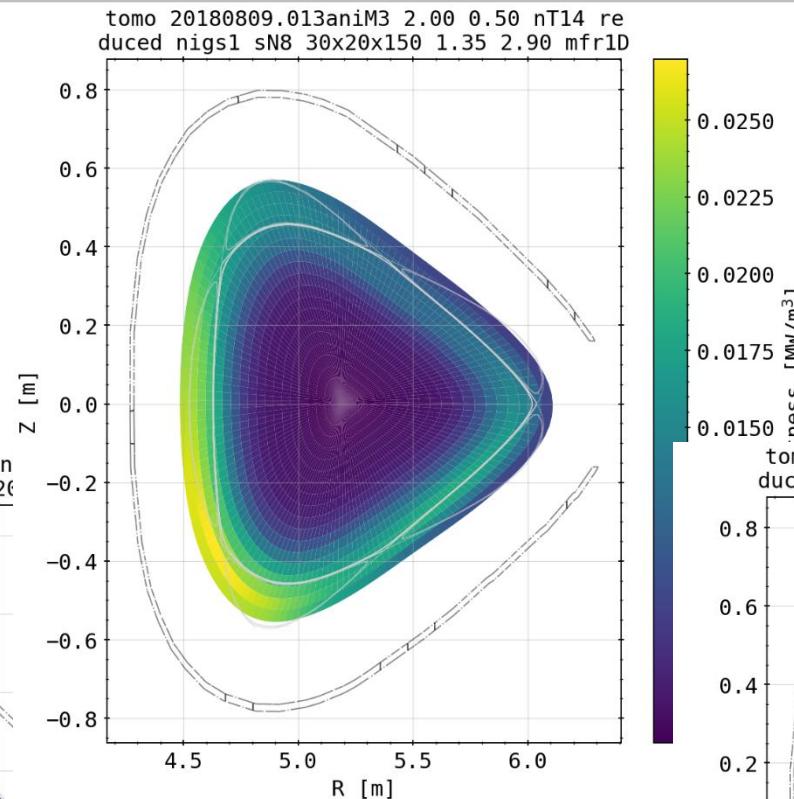
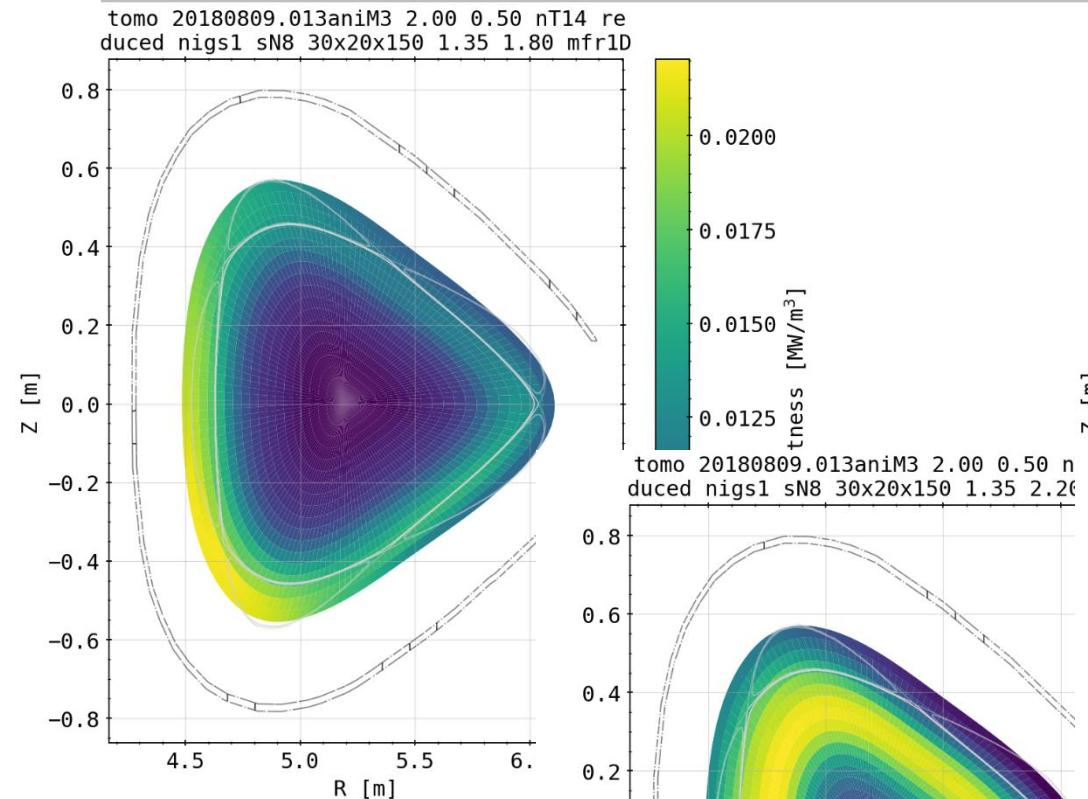
outside asymmetry,
assuming asymmetry



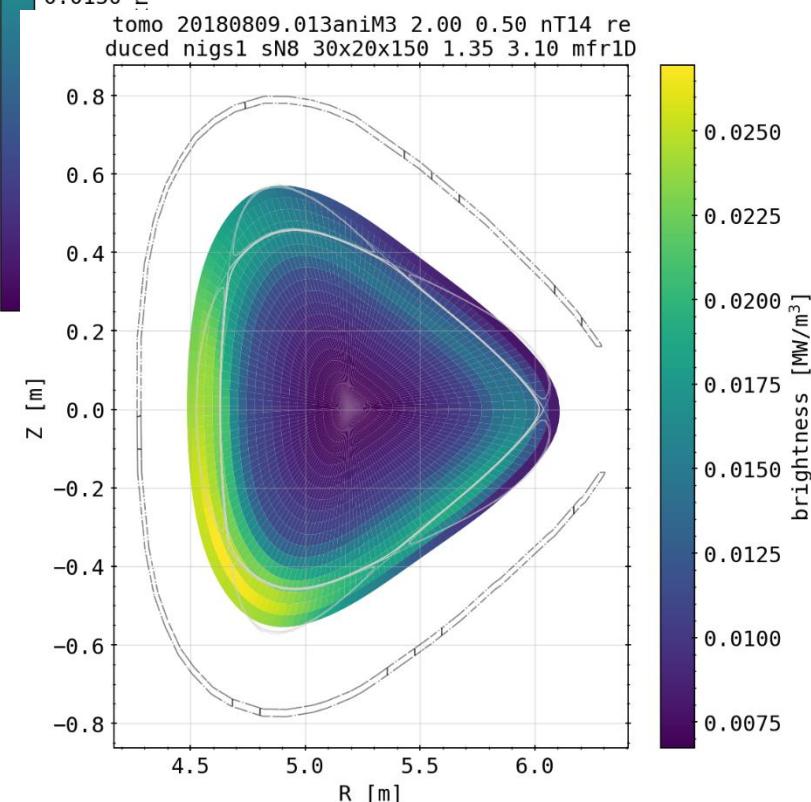
outside symmetry,
assuming asymmetry



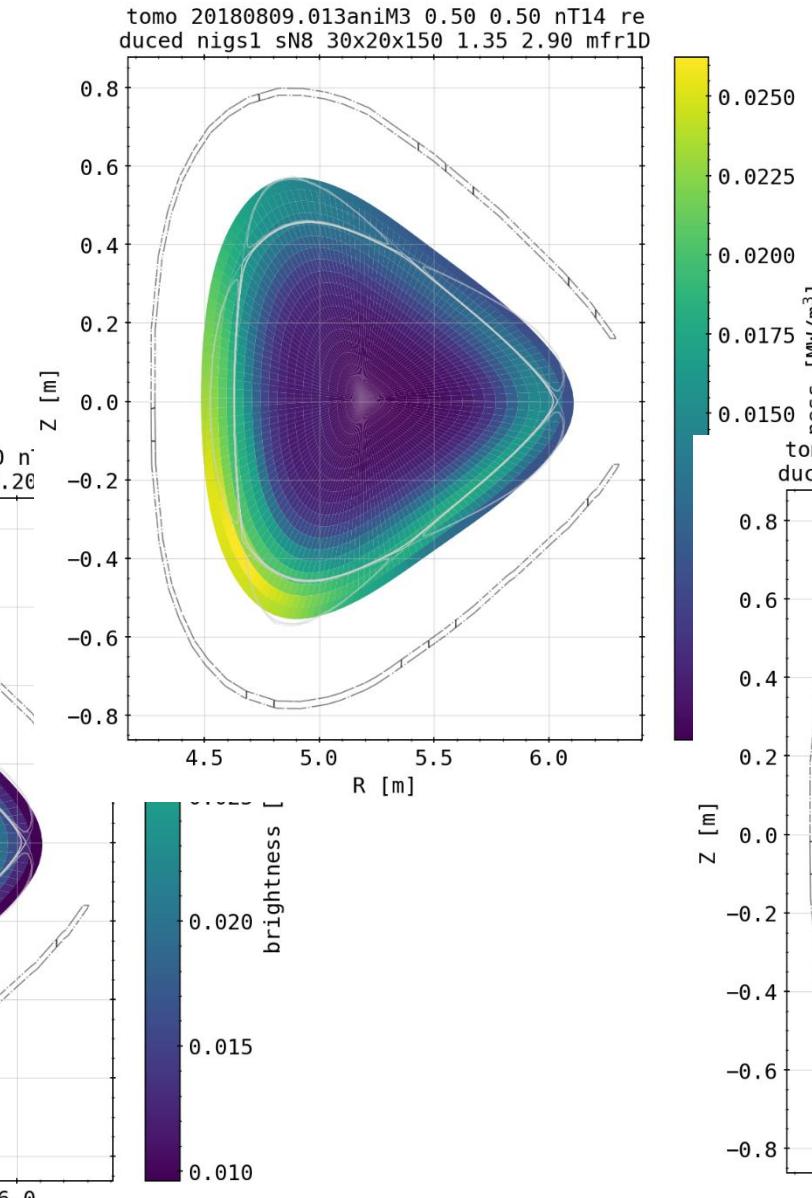
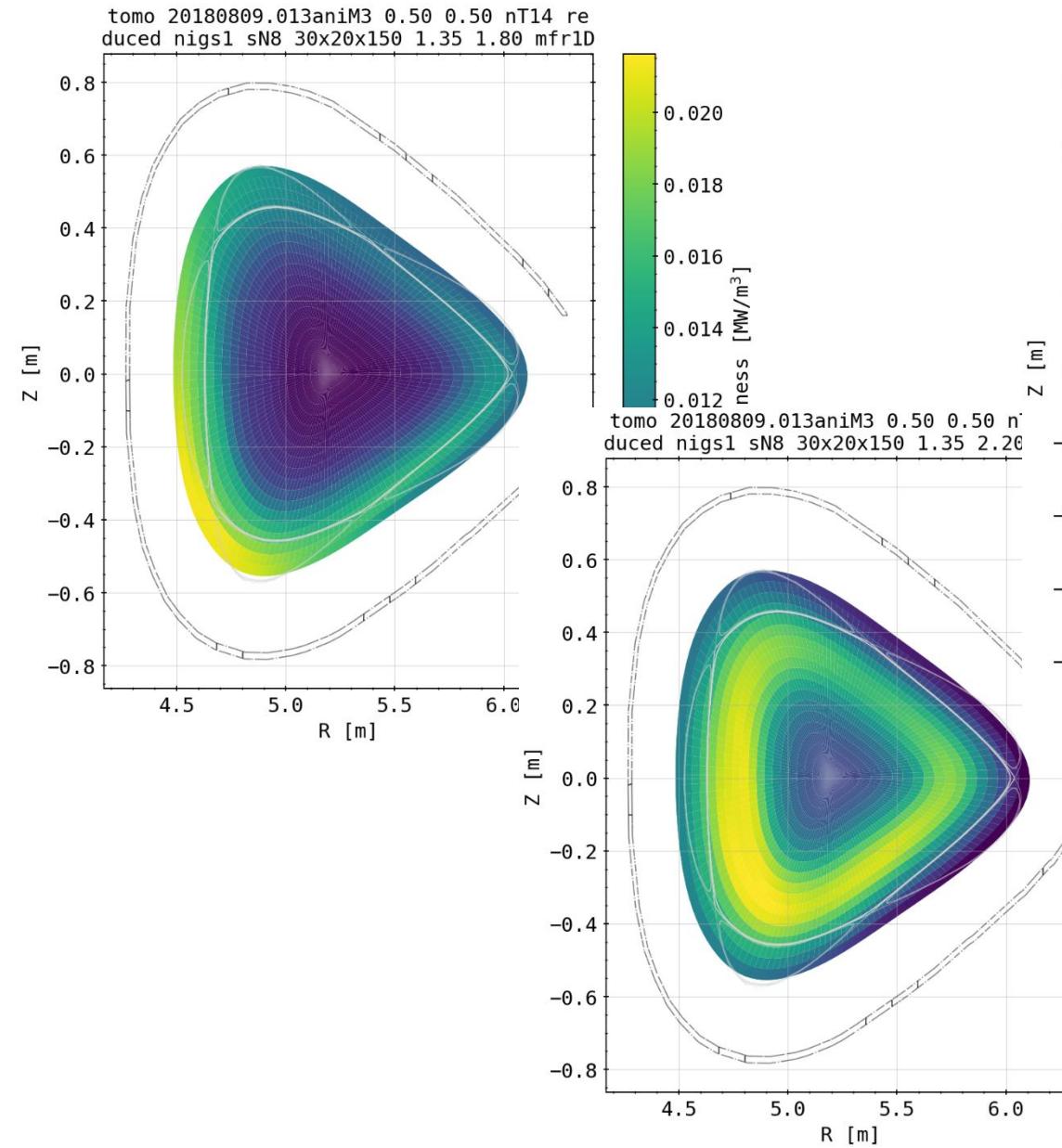
Asymmetry Propagation: 20180809.013



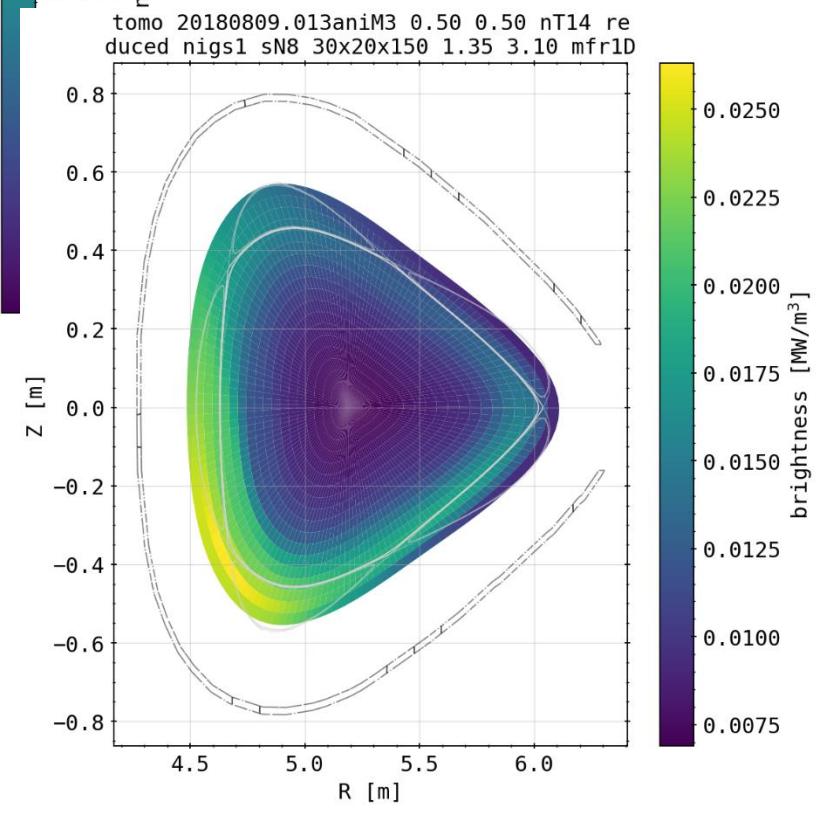
(core symmetric,
SOL asymmetric)



Asymmetry Propagation: 20180809.013

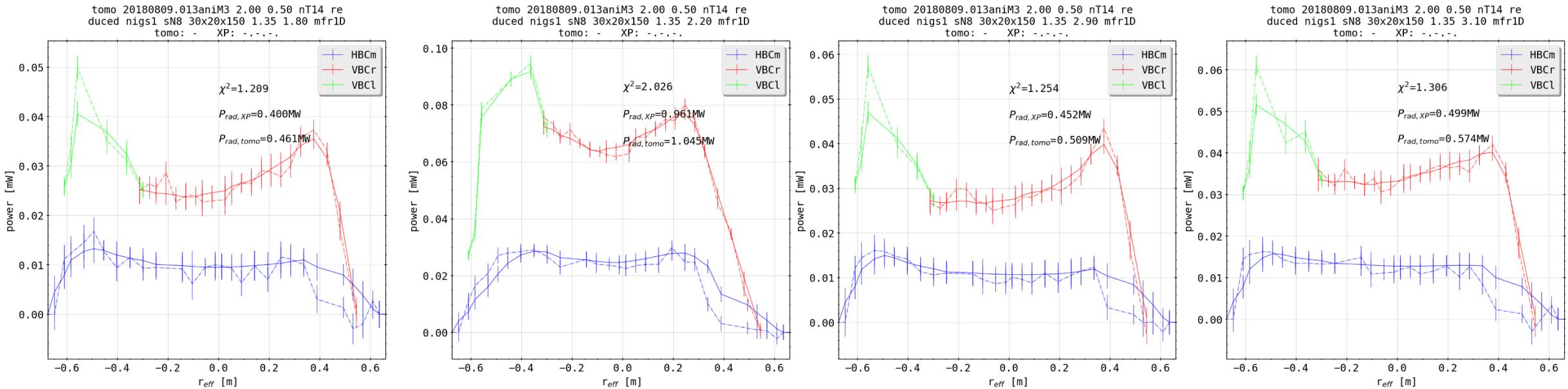


(core symmetric,
SOL asymmetric)



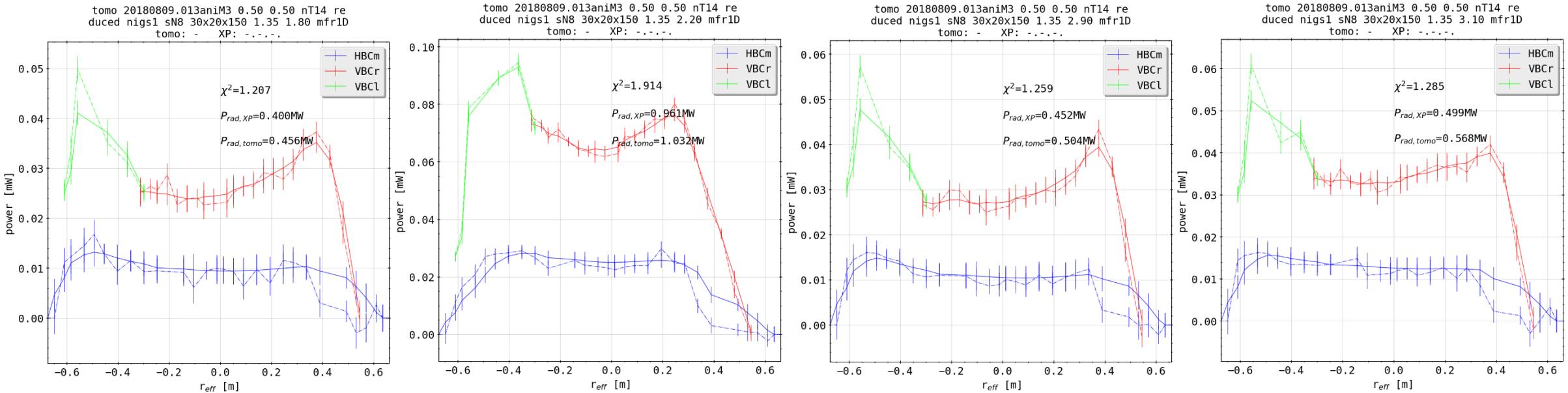
Asymmetry Propagation: 20180809.013

(core symmetric,
SOL asymmetric)

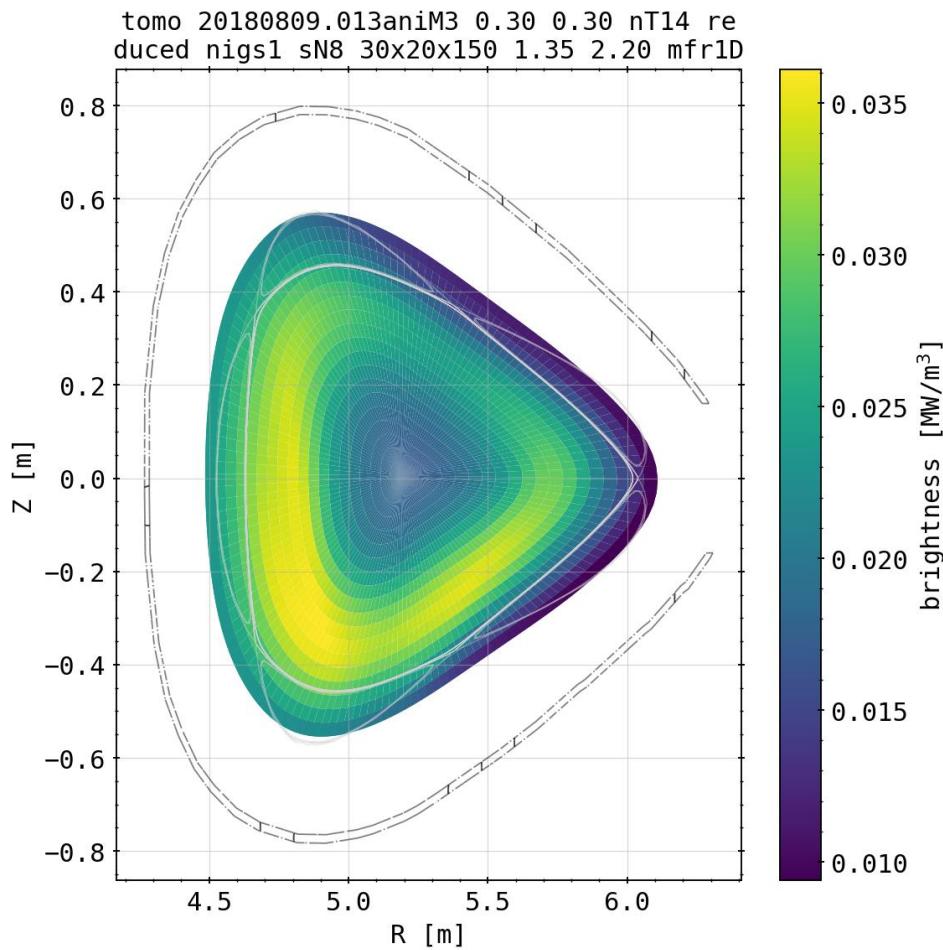


Asymmetry Propagation: 20180809.013

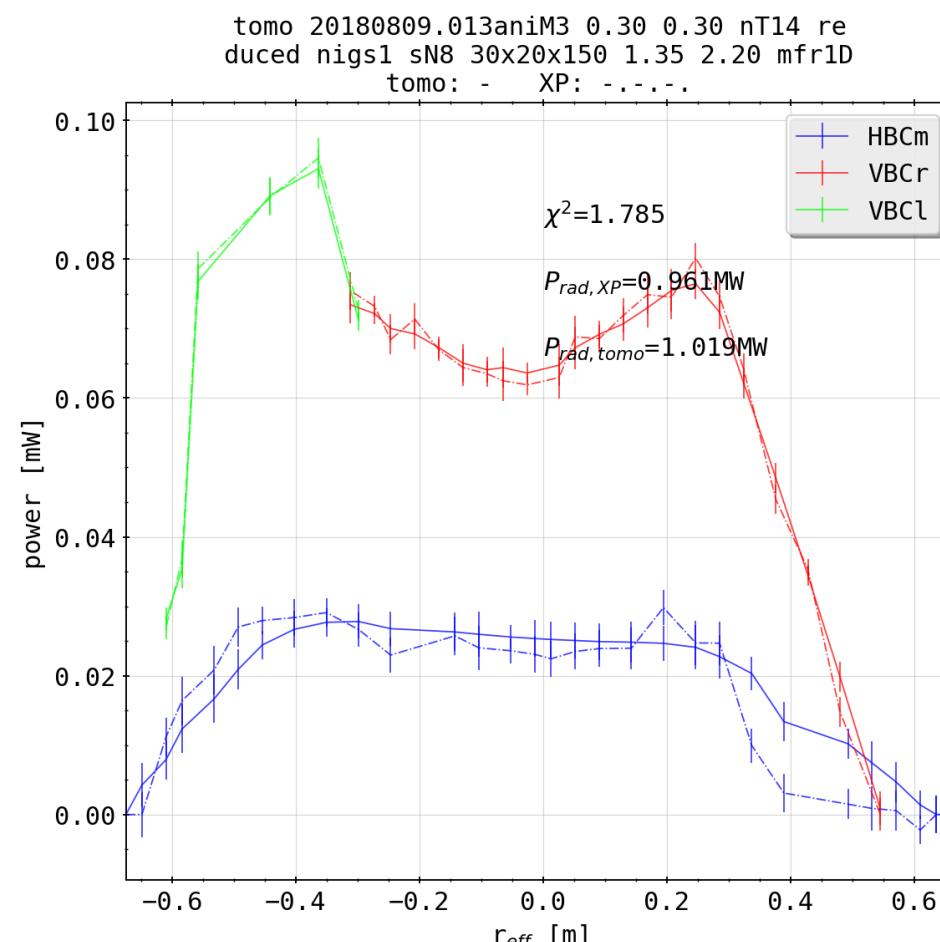
(core asymmetric,
SOL asymmetric)



Asymmetry Propagation: 20180809.013



(core asymmetric,
SOL asymmetric)



Impurity Seeding: Multi Chamber Model

two chamber model:

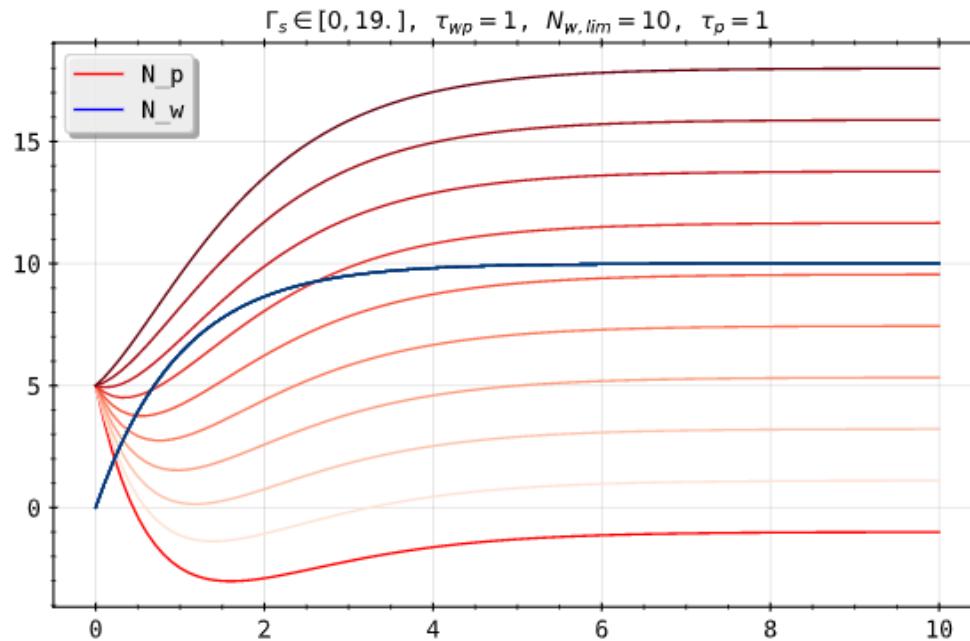
$$\dot{N}_p = \Gamma_s + N_w \tau_{w,p} - N_{w,lim} \tau_{w,p} N_p - N_p \tau_p$$



$$\dot{N}_w = (N_{w,lim} - N_w) \tau_{w,s}$$



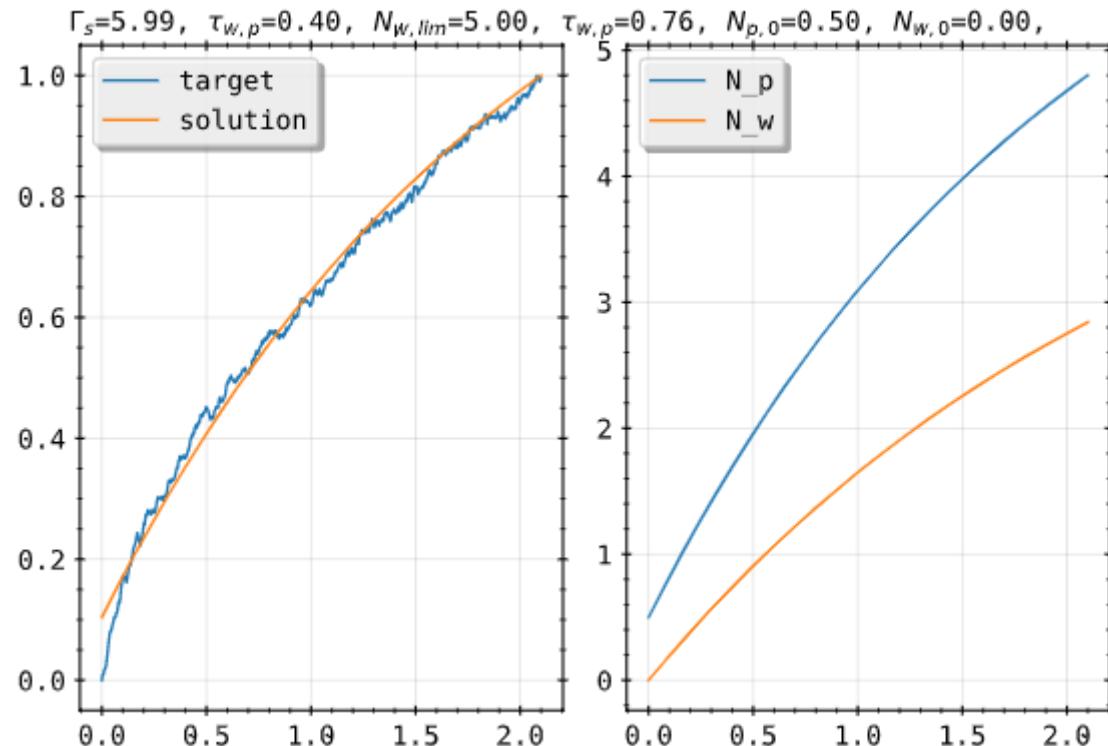
seeded pumping



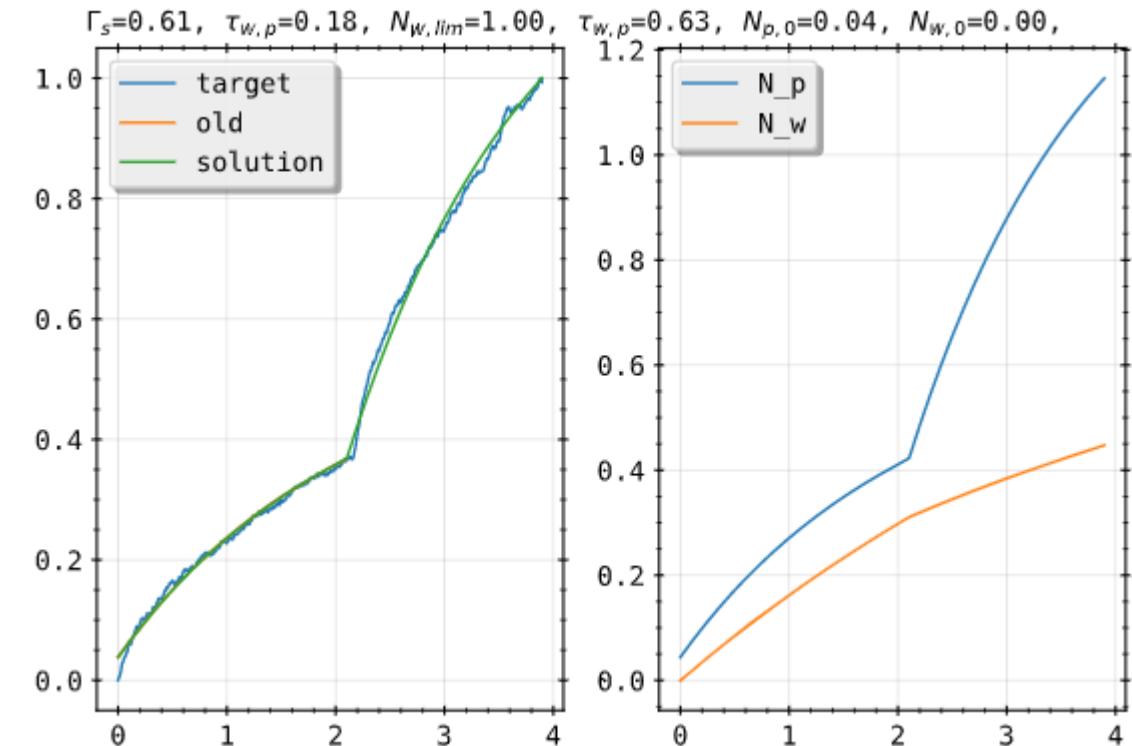
Impurity Seeding: Multi Chamber Model

two chamber model:

$$\dot{N}_p = \Gamma_s + N_w \tau_{w,p} - N_{w,lim} \tau_{w,p} N_p - N_p \tau_p$$



$$\dot{N}_w = (N_{w,lim} - N_w) \tau_{w,s}$$



Impurity Seeding: Multi Chamber Model

three chamber model:

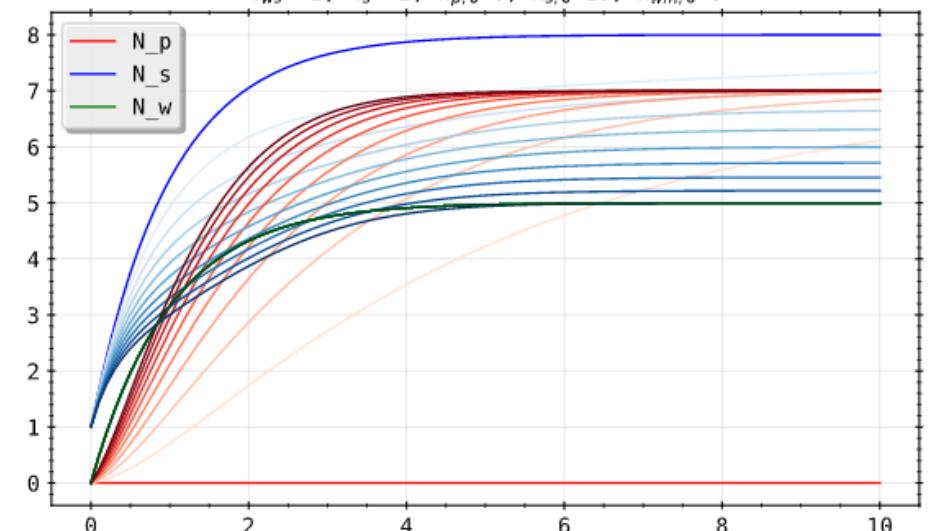
$$\dot{N}_s = \Gamma_s + \frac{N_p N_s \tau_{s,p}}{N_{p,lim}} - N_{w,lim} \tau_{w,s} - N_s (\tau_{s,p} + \tau_s)$$



$$\dot{N}_p = N_s \tau_{s,p} - \frac{N_p N_s \tau_{s,p}}{N_{p,lim} \tau_{p,s}}$$

$$\dot{N}_w = (N_{w,lim} - N_w) \tau_{w,s}$$

$$\begin{aligned}\tau_{sp} \in [0, 2], \quad N_{p,lim} = 10, \quad \tau_{ps} = .7, \quad \Gamma_s = 20, \quad N_{w,lim} = 5 \\ \tau_{ws} = 1, \quad \tau_s = 1, \quad N_{p,0} = 0, \quad N_{s,0} = 1, \quad N_{w,0} = 0\end{aligned}$$



Impurity Seeding: Multi Chamber Model

three chamber model:

$$\dot{N}_s = \Gamma_s + \frac{N_p N_s \tau_{s,p}}{N_{p,lim}} - N_{w,lim} \tau_{w,s} - N_s (\tau_{s,p} + \tau_s)$$

$$\dot{N}_p = N_s \tau_{s,p} - \frac{N_p N_s \tau_{s,p}}{N_{p,lim} \tau_{p,s}}$$

$$\dot{N}_w = (N_{w,lim} - N_w) \tau_{w,s}$$

$\tau_{s,p}=0.50, N_{p,lim}=0.61, \tau_{p,s}=13.35, \Gamma_s=80.49, N_{w,lim}=0.24, \tau_{w,s}=0.11,$
 $\tau_s=7.00, f=1.00, N_{p,0}=0.00, N_{s,0}=1.00, N_{w,0}=0.00,$

