



Max-Planck-Institut für Plasmaphysik

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Improvement on Reconstruction and Chordal Profiles





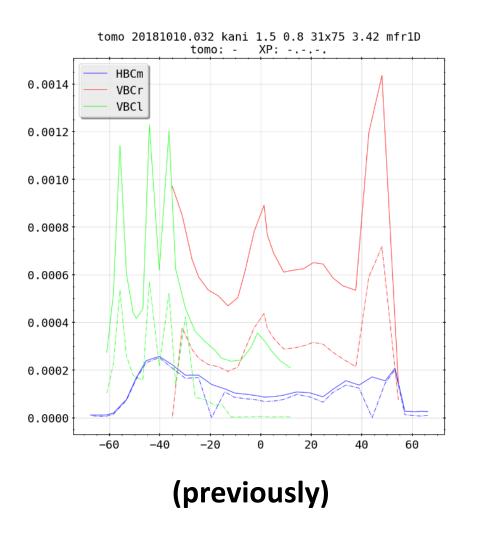
- Forward and backward calculation now using the emissivity matrices themselves, instead of channel geometries individually, hence calculating directly to and from the tomogram/phantom
- > comparing to input chordal profiles rather than archived ones
- ➤ also found error in channel selection and mapping that cause restrictions to zero where no data has been provided or excluded
- deleted broken and unusable channels from script

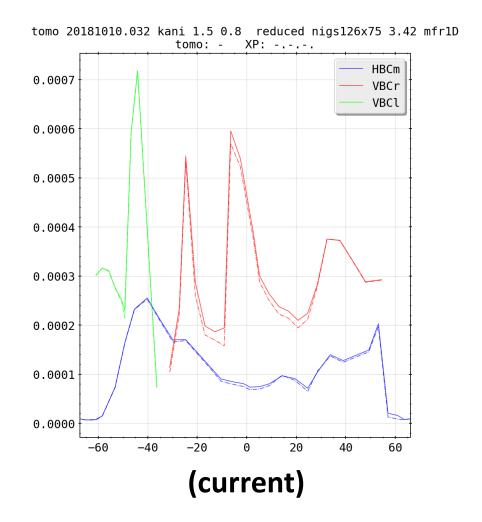
```
def forward integral(
    channels={'none': None},
    data=np.zeros((33 * 51)),
    emissivity=[np.zeros((32, 33 * 51))],
    debug=False):
chord = np.zeros((128))
for n, cam in enumerate(['HBCm', 'VBCl', 'VBCr']):
        nCh = channels['reduced']['eChannels'][cam]
        nCh = channels['full']['eChannels'][cam]
    tm2D = emissivity[n]
    forward = tm2D.dot(data)
        print(np.shape(forward), np.shape(nCh))
    for c, ch in enumerate(nCh):
            print('2', cam, c, ch)
        chord[ch] = forward[c]
return (chord)
```

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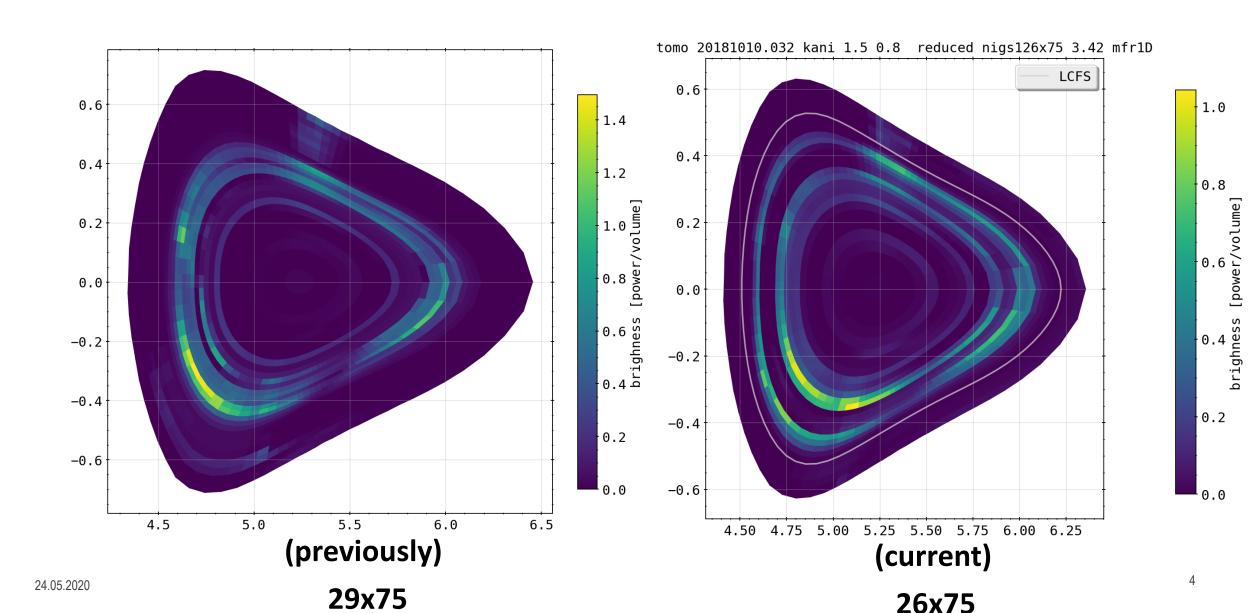




Improvement on Reconstruction and Chordal Profiles



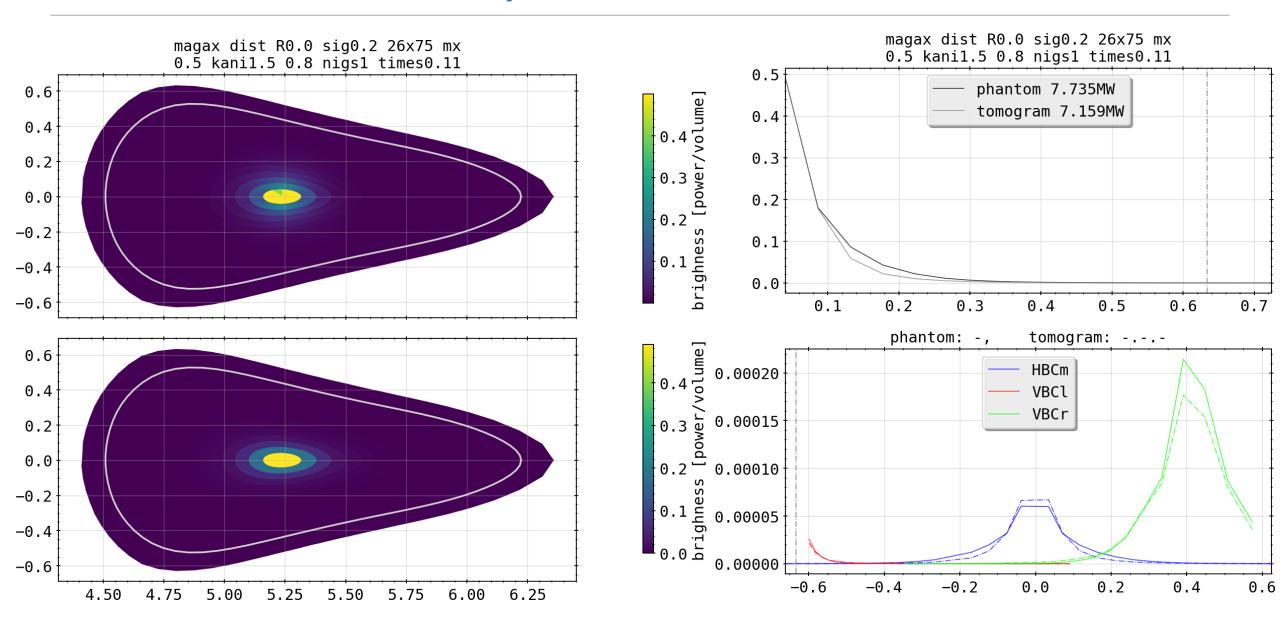




New Phantom Tests: Hot-Spot







Repositories





> personal routines and procedures specifically for calculation in

https://git.ipp-hgw.mpg.de/pih/QSB_Bolometry/-/tree/master/libinversion/libcalc

>private and/or locked git in

https://git.ipp-hgw.mpg.de/pih/bolometer_mfr

containing the MFR routines of D.Zhang, H.Thomsen

Repositories





> now reading and writing the radial profile of anisotropy factors, investigation to be done

First tests of X² for 2D profile done, needs better math (divergent if one profile zero in cell, normalization?); implementing D.Zhang's method?

>still need to look at actual MFR routine for k_ani exceptions and (if) possible cartesian test