

#### 2D genus topology of 21cm differential brightness temperature during cosmic reionization

Kyungjin Ahn (Chosun), **SEH** (CNU/KAIST), Changbom Park, Juhan Kim (KIAS), Ilian T. Iliev (Sussex) and Garrelt Mellema (Stockholm) arXiv:1008.3914

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### Redshifted 21cm signal







•  $g_{2D}(T_{th}) = (\# \text{ of hot spots}) - (\# \text{ of cold spots})$ Melott et al. 1989; Gott et al. 1990; Colley & Gott 2003; Gott et al. 2007



Simulations

N-body: GOTPM Dubinski et al. 2004; Kim et al. 2009

matter density

halo profile

matter density

halo profile

matter density

halo profile

Reionization: C<sup>2</sup>Ray Mellema et al. 2005 Mellema tal. 2005 differential brightness temperature

ionization fraction

ionization fraction

 $\delta T_b =$ 

differential brightness temperature

differential brightness temperature

$$\frac{(28 mK)\left(\frac{1+z}{10}\right)^{\frac{1}{2}}}{10}(1+\delta)(1-x)$$

time

#### Simulations

- N-body simulation
  - ACDM model with WMAP 5yr parameters
  - 2048<sup>3</sup> particles
  - $66 \text{ Mpc/h box} (\sim 30' \text{ at } z = 14)$
- Reionization simulation
  - 256<sup>3</sup> mesh
  - 4 source property models
    - 2 for high-mass halos only (M >  $10^9 M_{sun}$ )
    - 2 for high-mass and low-mass halos  $(10^8 < M/M_{sun} < 10^9)$

### Mock 21-cm sky map



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#### 2D genus: evolution process



2D genus: evolution process



2D genus: evolution process



#### 2D genus: evolution process





Low-mass source



- High-mass ( $M \ge 10^9 M_{\odot}$ ) source emissivity: same
- Low-mass  $(10^8 \le \frac{M}{M_{\odot}} \le 10^9)$  source emissivity:
  - f125\_125S: low
  - f125\_1000S: high





(a)

f125\_125S



f125\_1000S

high low-mass efficiency



more detailed bubbles



amplitude increases













f125\_1000S

high low-mass efficiency all overdense region ionized

genus = 0



### Beam shape dependency

#### Gaussian

#### **Compensated Gaussian**

Crude approximation of actual beam with lack of large scale signal



Beam shape dependency





Beam shape dependency





Sensitivities in SKA





- 2D genus curve clearly shows the evolution of the reionization process.
- 2D genus method can be used to discriminate between various scenarios.
- SKA will be able to produce data suitable for the 2D genus analysis, with
  - Integration: 100 ~ 1000 hours
  - Beam size: 2 ~ 3 arcminutes
  - Bandwidth: 1 ~ 2 MHz



# Thank you!

