

Molecular Dynamic Simulation on Boron Cluster Implantation for Shallow Junction Formation

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INTRODUCTION

Boron cluster ion implantation is a potential technology for shallow junction formation in integrated circuits manufacture [1]. By cluster implantation, space-charge blow up of the beam can be minimized and more B atoms are implanted at the same beam current. Another advantage of using clusters is that it generates more defects to reduce channeling effect [2]. With the technology becoming more practicable, capability of simulating it for shallow junction formation becomes necessary, which is still difficult for full MD method. A localized molecular dynamic method [3] for simulating cluster implantation at normal dose, e.g. 10^{15} cm^{-2} , aimed at microelectronics application, is presented in this paper.

MODEL

In our MD simulation, a moving simulation box [3], composing of $3 \times 3 \times 3$ unit cells (about 200 atoms), is applied to save computing time. Accurate geometry structures of boron clusters [4] are considered in the model by described accurate position of both B and H atoms as is illustrated in Fig. 1. In order to describe interactions between atoms in boron clusters, potential function in which the repulsive action term is ZBL potential function and attractive term has the form of SW [5] potential function is applied.

RESULTS

Simulation examples of $\text{B}_{10}\text{H}_{14}$ and $\text{B}_{18}\text{H}_{22}$ are performed. The snapshot of cluster implantation is shown in Fig. 2. Simulation results are shown in Fig. 3~6. It can be seen that range profiles from simulation agree well with the SIMS data. Specially, agreement of H profile is also shown for $\text{B}_{18}\text{H}_{22}$

implantation, where obvious deviation at near surface region exists maybe due to uncertainty of SIMS measurements at such shallow region. Simulation and SIMS data of B monomer implantation at equivalent energy and dose is also shown in Fig. 3 and 5 as comparison. It shows cluster implantation tends to generate shallower distribution profile. It is notable that with cluster model presented, the simulation can reproduce that difference well.

Fig.4, 6 also show that cluster implantation induces more Si interstitial defects than B monomer. These results agree with reference [2].

ACKNOWLEDGEMENT

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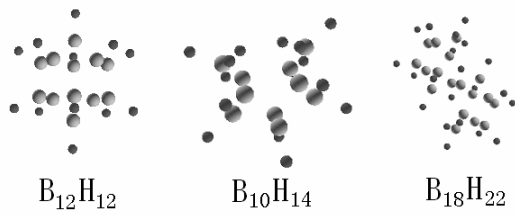


Fig. 1. The schematic representation of boron clusters ($B_{12}H_{12}$, $B_{10}H_{14}$ and $B_{18}H_{22}$). Structure of $B_{10}H_{14}$ and $B_{18}H_{22}$ is described based on that of $B_{12}H_{12}$.

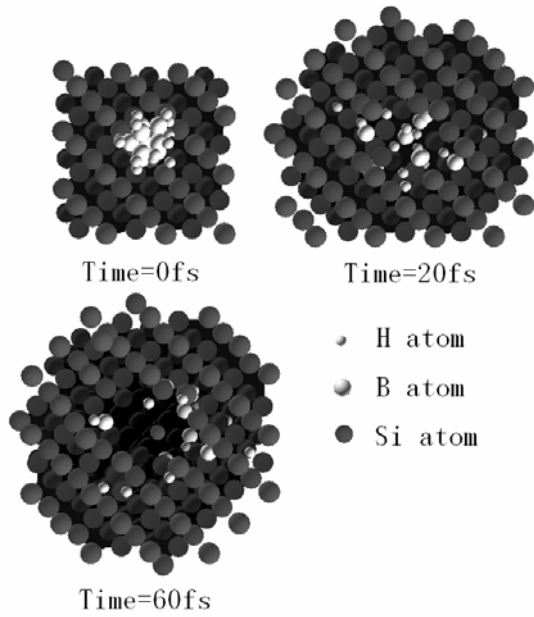


Fig. 2. Snapshots of $B_{10}H_{14}$ implantation at 1keV/molecular

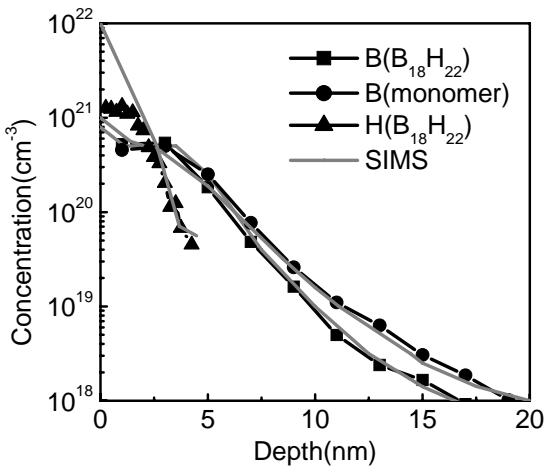


Fig.3 Concentration profiles of B atoms and H atoms for $B_{18}H_{22}$ implantation at 10keV/molecular and B monomer atom implantation at 0.5keV/atom. (Dose $3e14B/cm^2$)The SIMS data is from Ref. [6]

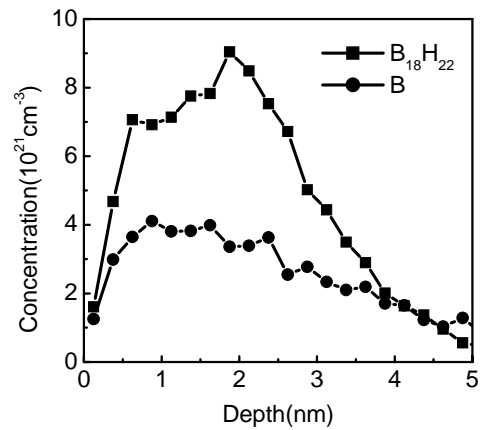


Fig.4 Concentration profiles of Si interstitials for $B_{18}H_{22}$ implantation at 10keV/molecular and B monomer atom implantation at 0.5keV/atom. (Dose $3e14B/cm^2$)

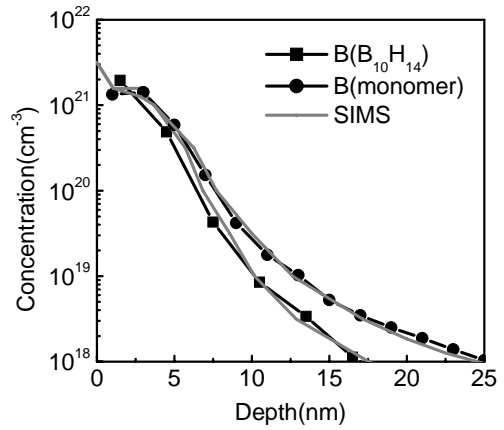


Fig.5 Concentration profiles of B atoms for $B_{10}H_{14}$ implantation (Dose $8.3e14B/cm^2$) at 5keV/molecular and B monomer atom implantation at 0.5keV/atom (Dose $8.1e14B/cm^2$). The SIMS data is from Ref. [7]

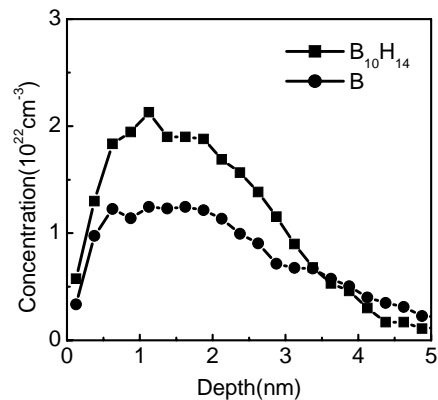


Fig.6 Concentration profiles of Si interstitials for $B_{10}H_{14}$ implantation (Dose $8.3e14B/cm^2$) at 5keV/molecular and B monomer atom implantation at 0.5keV/atom (Dose $8.1e14B/cm^2$).