GaCl will be shown. The analysis provides us useful information on the growth rate and the deposition composition. Next, HVPE growth of AIN and AlGaN will be described.

[1] Y. Kumagai et al. Phys. Stat. Sol. (c) 0 (2003) 2498.

[2] A. Koukitu at al. J. Cryst. Growth 281 (2005) 47.

Keywords: HVPE, AI-related nitrides, thermodynamic analysis

#### MS.49.3

Acta Cryst. (2008). A64, C88

# Fabrication of InN dot structures by droplet epitaxy using NH<sub>3</sub>

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Quantum dots (QDs) of nitride semiconductors have been studied intensively because of their usefulness for device applications. Recently, we have fabricated GaN dots by droplet epitaxy. The droplet epitaxy requires neither any anti-surfactants nor the induction of surface strain for the formation of dot structures, which means the possibility of the QD formation on any substrates. In this study, we investigated fabrication of InN dots by droplet epitaxy with changing substrate temperature. After nitridation of Si(111) substrates, 1 ML of In was deposited on the substrates at 200 °C , leading to the formation of In droplets. Then, NH3 gas was irradiated on the In droplets for nitriding. AFM observation showed that the average density of dots of  $1.1 \times 10^{10}$  cm<sup>-2</sup> at 450°C decreased as the temperature increased. Then, the density increased again up to  $4.0 \times 10^{10}$  cm<sup>-2</sup> at 570°C. However, no InN dot was observed on the sample nitridated

at 600  $^{\circ}$ C. In 3d XPS spectra in Fig.1 indicates that In-N component was increased as the temperature increased while In-O component was decreased. Almost all In atoms was found to bind to N atoms at 570  $^{\circ}$ C by the formation of InN crystalline.



Fig. 1 In 3d XPS spectra of InN dots grown by droplet epitaxy using NH3

Keywords: droplet epitaxy, indium nitiride, nitiriding

### MS.49.4

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# Dislocation density in silicon ingot during a unidirectional solidification process

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and dislocation density were also studied. The results reveal that long solidification time can reduce thermal stresses and dislocation density in a silicon ingot during the solidification process.



Keywords: solidification, stress, dislocations

### MS.49.5

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# AlN and GaN hetero epitaxy on Si substrate using activity modulation migration enhanced MBE

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In order to ontrol chemical and physical nitrogen activities two discharge modes of E and H modes[1] under a radio frequency inductive coupled (RF-ICP) discharge of nitrogen gas was investigated changing RF-power of nitrogen pressure. The chemical activities of disociated nitrogen atoms N+N\* are much higher than those of excited nitrogen molecules N2\*. However N2\*has small chemical activity but large physical activity, which has large energy to enhance migration of Al or Ga atoms on a growing surface. The E and H modes of the ICP discharge are low brightness (LB) and high brightness (HB) discharges, respectively. Dissociated atomic nitrogen species are created by only the HB mode. The time sequence of the opening of the shutter of a Ga effusion cell was used for the trigger signal to operate the mode change operation of nitrogen ICP -RF discharge. An activity modulation migration enhanced epitaxial growth (AM-MEE) method is prpposed as an application of the MEE[2]. In order to grow the high quality layers on a Si substrate the AM-MEE growth method was performed. The sequence of the LB and HB discharges during the AM-MEE the nitrogen flux irradiation plays an important role to control the chemical activity of nitrogen. Activity control of nitrogen is also applied to a fabrication method of beta-Si3N4 epitaxial layer on Si(111) as an initial buffer layer for AIN film.

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Phy 27 (1988) 169.

Keywords: activity, MBE, gallium nitrides

### MS.49.6

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# Contrast of dislocations in 4H-SiC by SR topography in grazing-incidence geometry

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Dislocations near surface of 4H-SiC were observed using synchrotron radiation topography in the Bragg case with grazingincidence geometry. Figure is an image of basal-plane dislocation half-loop at g=11-28,  $\lambda$ =0.15nm on Si-face. The (0001) plane is tilted towards the [-1-120] direction by 8 degrees from the surface. In this condition, lattice defects within 10 µm depth are observed. Along this dislocation line, bright contrast at A, dark and bright asymmetric line at B, and dark contrast at C are observed. Absence in contrast can be seen at B at g=1-108, and so that B is a screw dislocation part. We have observed migrations of dark dislocations in specimens after forward-bias degradation effect, in which Si-core dislocations are known to move. Thus we concluded that C is Si-core, A is C-core edge dislocation, and the Burgers vector is 1/3[-1-120]. The observed dark and bright contrast is discussed to be similar effect

described by Ando and Kato (1970). By applying this rule we could identify uniquely 6 different Burgers vectors for all basal-plane dislocations and threading edge dislocations at only one diffraction condition. Ando and Kato: J. Appl. Cryst. **3** (1970) 74.



Keywords: wide-bandgap semiconductors, dislocations, topography X-ray

### MS.50.1

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# Decoding homophilic recognition specificity of Dscam, a neuronal receptor with thousands isoforms

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The Dscam gene gives rise to thousands of diverse cell surface receptors thought to provide homophilic and heterophilic recognition specificity for neural development and immune responses. Mutually exclusive splicing allows for the generation of sequence variability in three immunoglobulin (Ig) ecto-domains (D2, D3, D7). X-ray structures of the N-terminal four Ig domains (D1-D4) of four distinct Dscam isoforms have been determined. The structures

reveal a horseshoe configuration, with variable residues of D2 and D3 constituting two independent surface-epitopes on either side of the receptor. All these four isoforms engage in homo-dimerization coupling variable domains D2 with D2 and D3 with D3 using the same epitope. The recognition specificity has been analyzed to decode how sequence and local conformation of these two variable domains contribute to homophilic interaction. The structure of the third Ig-like domain D7 has also been determined in the form of D7-D8 fragment for several isoforms. A general view of how these variable Ig domains embedded in thousands receptor isoforms offer hemophilic recognition for neuronal wiring has been provided.

Keywords: dscam receptor, decoding recognition specificity, thousand isoforms

### MS.50.2

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# Crystal structure of the [2Fe-2S] transcriptional activator SoxR bound to DNA

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SoxR functions as a sensor of oxidative stress such as superoxide and nitric oxide. It exists as a dimer with each subunit containing a [2Fe-2S] cluster. Reversible oxidation of the [2Fe-2S] cluster activates SoxR to enhance the production of various antioxidant proteins through the soxRS regulon. SoxR belongs to the MerR family of transcriptional activators, target promoters of which have an unusual 19 or 20 bp spacer between the -35 and -10 operator elements. In the active state, SoxR and other MerR family proteins activate transcription from unique promoters by distorting the DNA conformation. In order to elucidate structural features of the ironsulfur cluster of SoxR and the transcriptional activation mechanism, we have determined the crystal structures of SoxR and its complex with DNA in the oxidized (active) state [1]. The overall structure of SoxR consists of a DNA binding domain, a dimerization helix and an Fe-S cluster binding domain. The dimerization helix forms an antiparallel coiled-coil, stabilizing the SoxR dimer. The structures reveal that the [2Fe-2S] cluster of SoxR is unusually solvent-exposed and surrounded by an asymmetric environment, suggesting that the asymmetrically charged environment is a key factor of redoxdependent conformational changes of SoxR and the target promoter. The DNA structure is shown to be sharply bent at the middle and unwound by 3-bp, compared to a B-form DNA. Based on comparison of the target promoter sequences of the MerR family, the present structures shows an activated promoter conformation with a 20-bp spacer in the MerR family.

[1] Watanabe S, Kita A, Kobayashi K, Miki K., Proc Natl Acad Sci USA, 2008, 105, 4121.

Keywords: SoxR protein, MerR family, transcription factors

### MS.50.3

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Hybrid LRR technique and crystal structures of the toll-like receptor complexes

Jie-Oh Lee