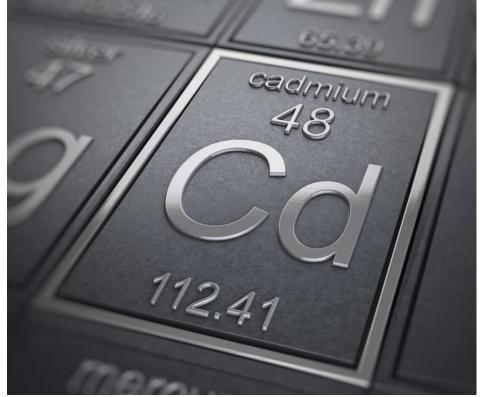


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Genetic variation as a tool for Cd mitigation

- Grafting high Cd varieties on low Cd varieties has been shown to reduce Cd accumulation in the shoots of the high Cd variety (Chao et al. 2012; Sugiyama et al. 2007)
- Proteins involved in Cd accumulation have been identified HMA 2, 3, 4;
 Nramp 1, 5, 6; PCS
- Breeding programmes that target loss of function genes (e.g. mutant screening, TILLING) (Chen and Ma 2016)
- Work mainly done in Arabidopsis, <u>rice</u>, Noccaea caerulescens (pennycress)

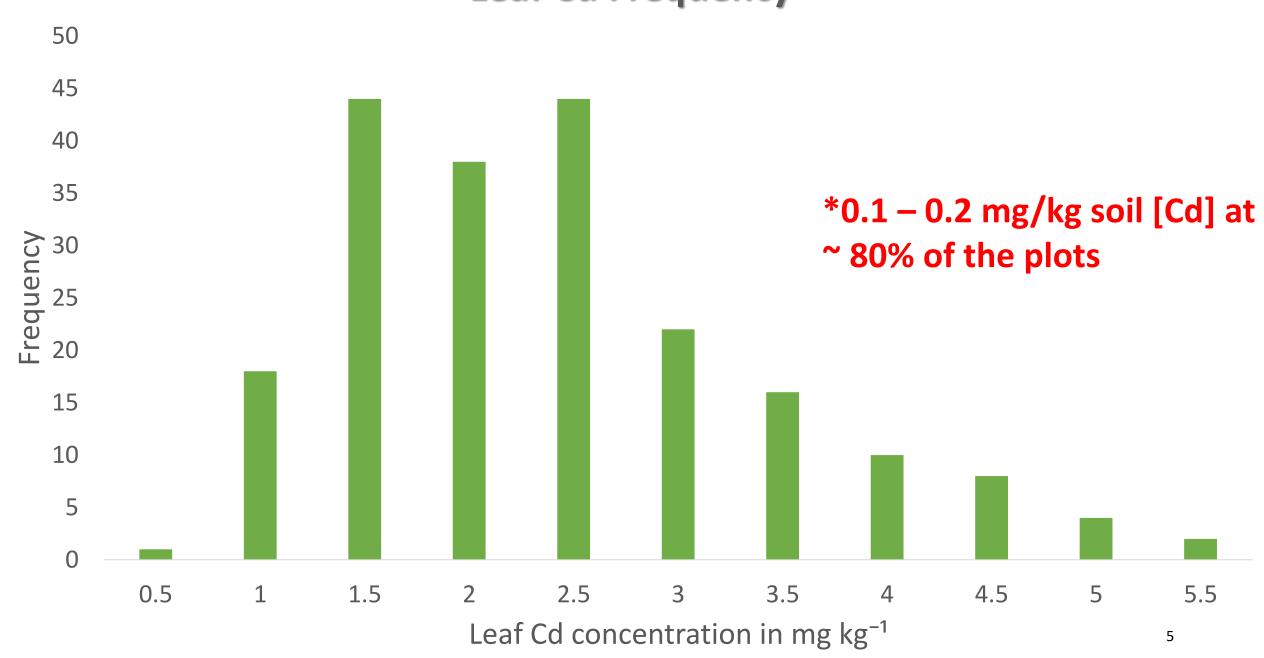
Gap

 The mechanism of Cd uptake and partitioning within the cocoa plant is unknown.

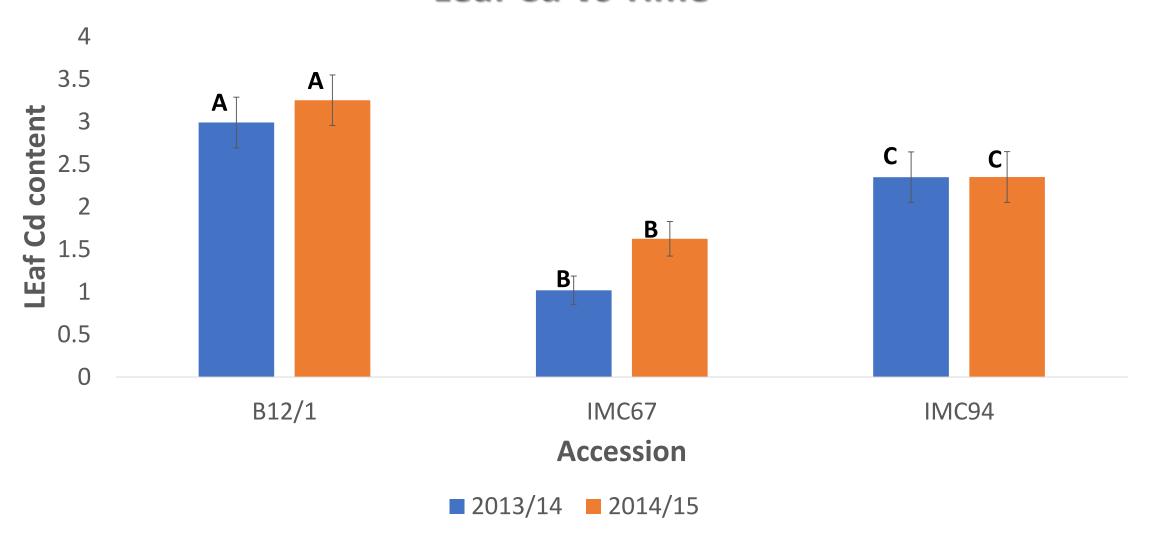


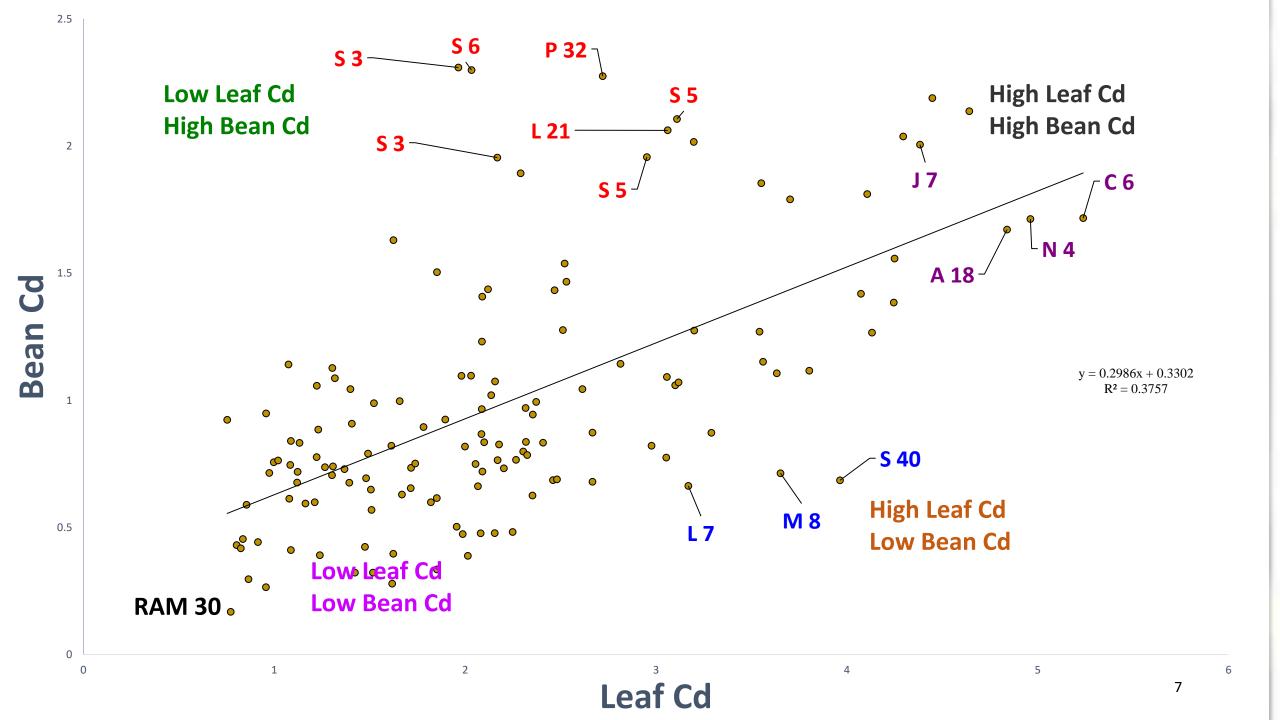


Leaf Cd Frequency

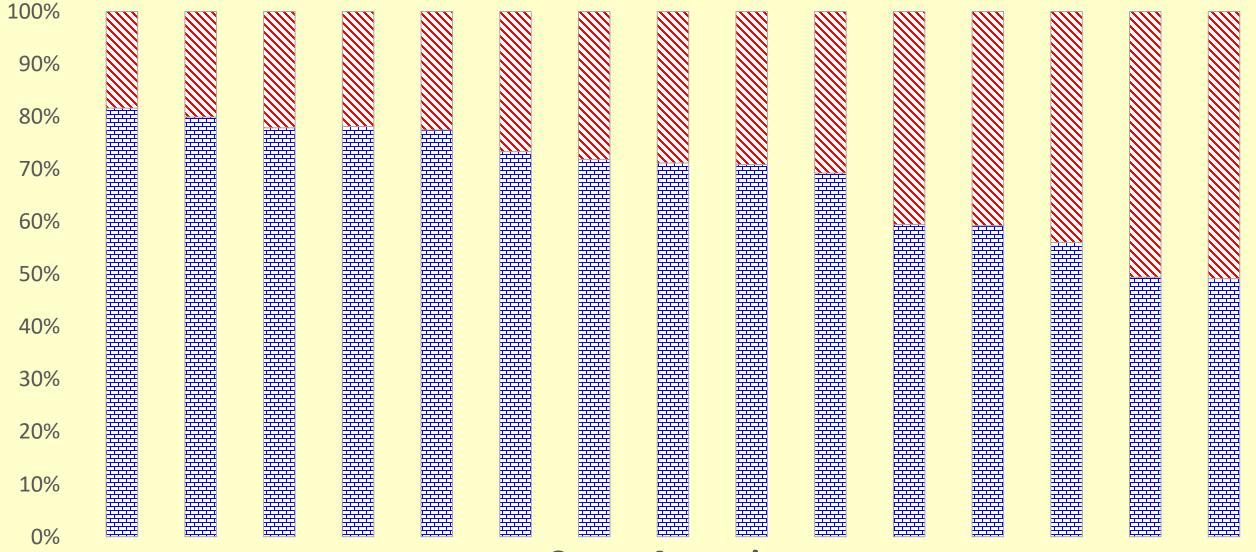


Leaf Cd Vs Time



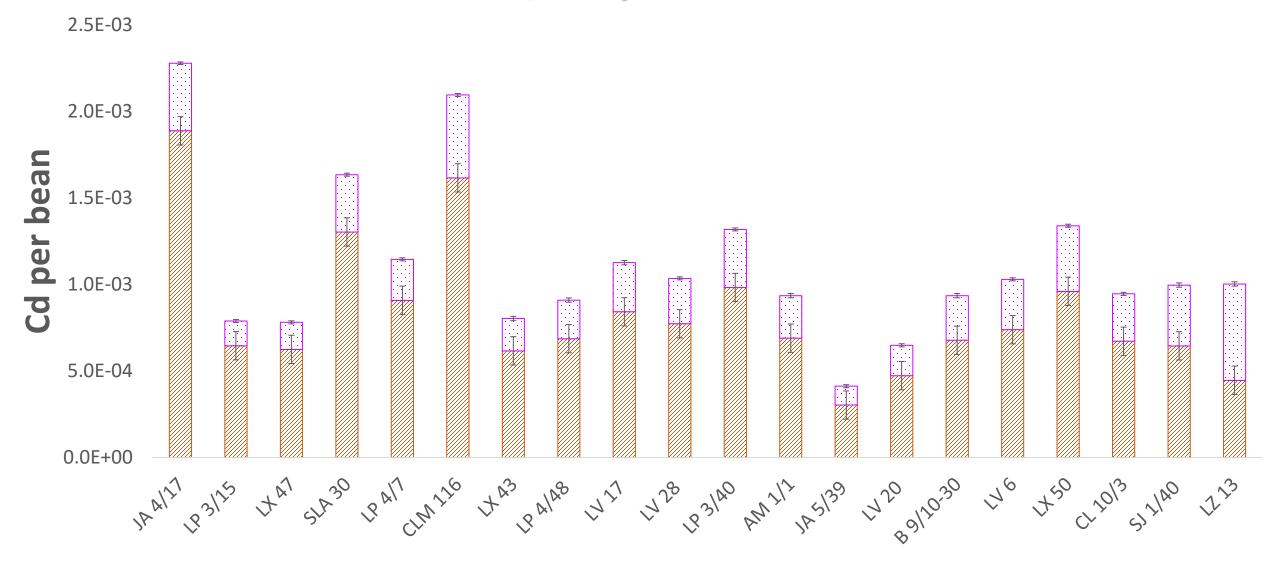


Bean/Leaf Cd Ratio



Cocoa Accession

Testa/ Cotyledon Ratio



Accession

Conclusions

- ICGT is the ideal site to screen for genetic variation in Cd bioaccumulation
- Variation of leaf and bean Cd between accessions probably due to genetic variation
- Multiple regulatory points during the journey of Cd from the soil to the cotyledon

Future Work

 Study of a larger number of accessions to identify the major pathways of Cd accumulation in cocoa

Acknowledgements

Financial support was provided by the ECA/ Caobisco/ FCC Joint Cocoa Research Fund

