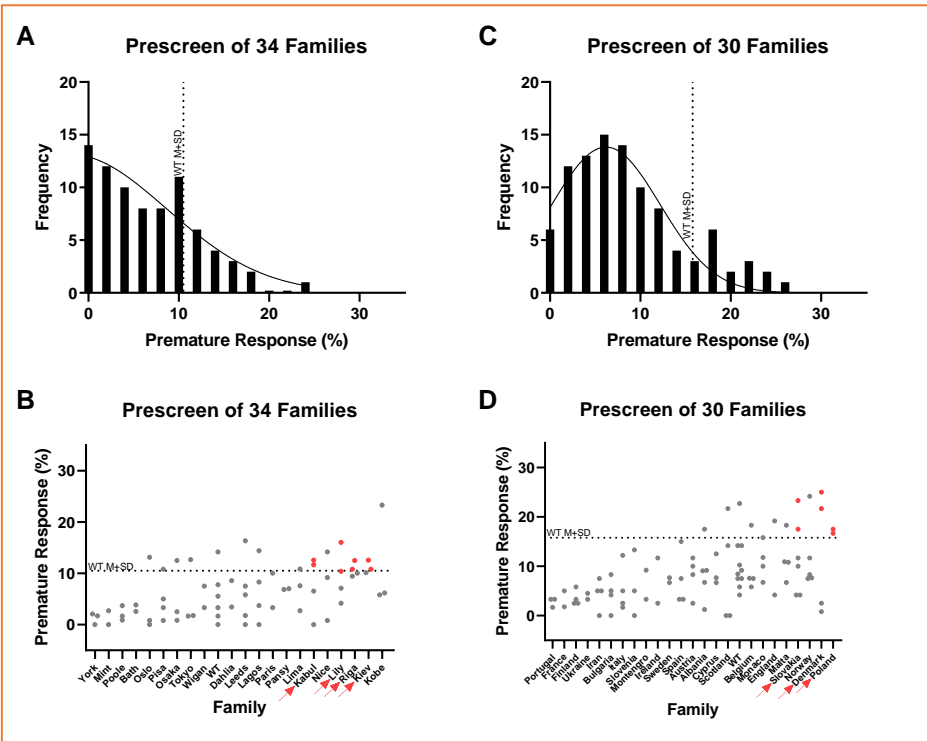


# A 5-Choice Serial Reaction Time Task Screen of ENU-Mutagenised Zebrafish Identifies Lines Showing Heritable Impulsivity and Two Mutations Cosegregating with Impulsivity

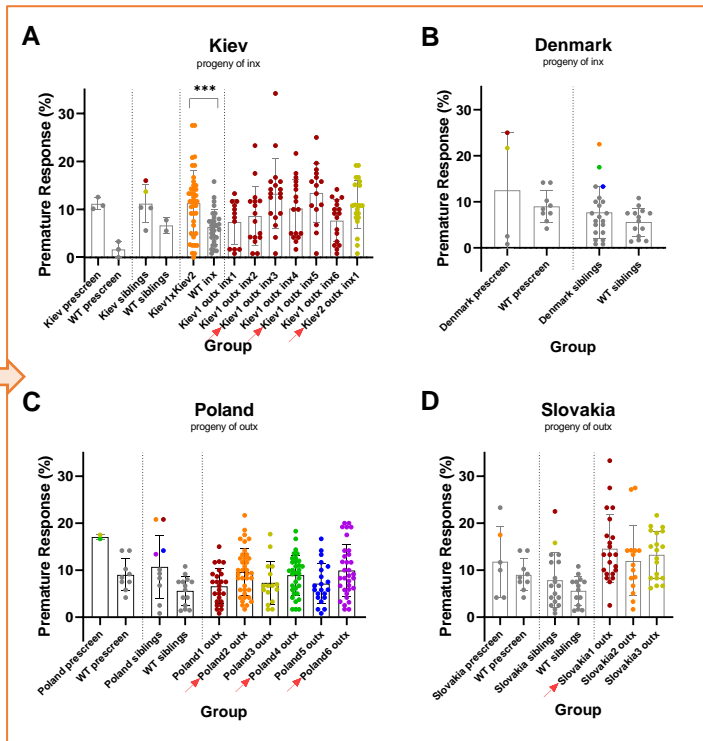
Saeedeh Hosseinian, Will Havelange, Munise Merteroglu, Adele Leggeri, Aleksandra Mech, Sofia Angianni, Ian Sealy, Elisabeth Busch-Nentwich, Caroline Brennan

School of Biological and Behavioural Sciences, Queen Mary University of London, London E1 4NS

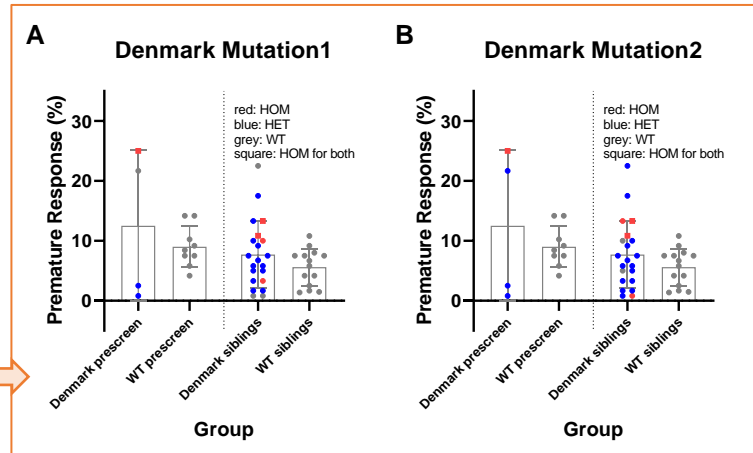


We aimed to identify genes and pathways underlying impulsivity as a risk factor for addiction.

- 1) We generated 102 F3 ENU families covering 7926 dominant and 3554 recessive alleles.
- 2) Prescreen of 6 fish from 102 families identified 7 candidate families with at least 2 fish showing premature response higher than WT M+SD.



- 3) Siblings of 4 candidate families were screened and confirmed the phenotype.
- 4) Next generation of 3 candidate families were screened and confirmed the heritability of the phenotype.



- 5) Exome sequencing of F1 of 7 candidate families identified 113 candidate mutations of which 10 are associated with impulse control disorders in GWASs.
- 6) Competitive allele specific PCR of F3 of 3 candidate families identified 2 candidate mutations in a candidate family cosegregating with the phenotype.
- 7) Confirmation and characterisation of ENU and CRISPR-Cas9 lines are ongoing.

This is the first forward genetic screen for impulsivity showing potential success.