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Gene-environment interactions involve DNA methylation from early development to adult life, and maybe over generations.

The study of genetic variations together with epigenetics may reveal higher resolution genomic association pictures.

Study of RA and MS revealed genetic/epigenetic associations, new risk/protective genotypes in MS and genotype/ epigenotype interactions with smoking in the risk risk for disease.

Smoking affects DNA methylation and hydroxymethylation of specific genes in bronchoalveolar lavage cells (macrophages). May be crucially important to separate 5-mC from 5-hmC.

The chronic alcoholic brain demonstrates genotype dependent DNA methylation close to the *DLGAP2* gene, whose product is implicated in glutamatergic transmission. Therapy for alcohol dependence may involve the newly found target.



Acknowledgements

Malin Almgren Louise Sjöholm Mikael Ringh Joëlle Rüegg

Maja Jagodic Lara Kular Francesco Marabita

Lars KLareskog Leonid Padyokov

Georgy Bakalkin, Uppsala Daniil Sarkisyan, Uppsala

Karolinska

Andy Feinberg, Johns Hopkins Yun Liu, Fudan Victor Karpyak, Mayo