



## 7<sup>th</sup> EpiCoN meeting Program



June 12<sup>th</sup>, 2018 – Astrid Fagraeus A103

09:00 – 09:15 Welcome and announcement of the program.

09:15 – 10:20 Session 1 – Do we care about H3K36 methylation?

09:15 – 09:45 Isaac Dorafshan – Methylation of H3K36 is irrelevant for anti-Polycomb function of Ash1

09:50 – 10:20 Henrik Lindehell – Chromosome-wide transcriptional responses upon disturbed H3K36 methylation machinery

10:25 – 11:30 Let's discuss!!!

11:30 – 12:30 Lunch break

12:30 – 14:15 Session 2 – Involvement of DNA and RNA modifications in cancer

12:30 – 13:00 Emma Andersson-Evelönn – CpG methylation can predict outcome in ccRCC

13:05 – 13:35 Zahra Haider – DNA methylation subgroups in T-cell Malignancies

13:40 – 14:10 Cyrinne Achour – Understanding the role of the m<sup>6</sup>A methyltransferase METTL3 in breast cancer

14:15 – 14:30 Coffee break

14:30 – 15:35 Session 3 – Genetics meets epigenetics

14:30 – 15:00 Tatiana Kan – DNA elements involved in Polycomb regulation of CCND2 transcription

15:05 – 15:35 Juan Barrasa – Targeting of PRC1 to human genome

15:40 – 16:00 Time for questions and conclusions / Closing Remarks

17:00 – 18:30 Fun activity (Mariehemsångarna)

18:30 – 21:00 Dinner

