



Talks will be 10 mins with 5 mins for questions

8:00 Clare Turnbull: Saturation genome editing of *BAP1* and the implementation of these data in clinical practice

8:15 Catherine Zhou: Dutch Early-Stage Melanoma (D-ESMEL) study

8:30 Thamila Kerkour: Who is at risk for distant metastasis of early-stage melanoma? Results of genomics analyses in the D-ESMEL study

8:45 Veronica Höiom: Polygenic Risk Scores for Melanoma in the Swedish population

9:00 Daniela Robles Espinoza: Ancestry and somatic profile in melanoma

9:15 Anne Cust: Melanoma risk assessment in practice

9:30 Aideen McInerney-Leo Mainstreaming melanoma genetic testing

9:45 Wrap-up and Discussion