



Functional Annotation of all Salmonid Genomes (FAASG)

FAASG Satellite Meeting to ICISB 2019.

Nov 21st 2019, Royal College of Surgeons, Edinburgh

0830 - 0900: *Coffee and registration*

0900 - 0930: **Welcome, intro to FAASG, scene setting**

0930 - 1030: **Salmonid reference genomes – where are we now & where do we need to get?**

- CIGENE lab – scene setting and TRANSPOSE project (20 mins)
- Koop lab - Pacific salmon spp. (10 mins)
- Palti lab – USDA trout/salmon (10 mins)
- 5 min slots available for relevant updates (on request)
- Group discussion (remaining time)

1030 - 1100: *Coffee*

1100 - 1230: **Progress since the whitepaper: funded functional annotation projects, links with FAANG, links with Ensembl**

- Scene setting (5 mins)
- ASA working group: Update on progress (15 mins)
- BDA working group: Update on progress (15 mins)
- MDS working group: Update on progress (5-10 mins)
- PHE working group: Update on progress (5-10 mins)
- Short slots to present FAASG-relevant projects or progress (open to all)
 - USDA rainbow trout project
 - AQUA-FAANG & H2020 projects
 - Yniv Palti – Feedback on FAANG-Mine
 - Open 5 min slots (on request)
- Group discussion (remaining time)

1230 – 1300: Role of EMBL-EBI in supporting FAASG

- Peter Harrison, FAANG data coordination centre (15-20 mins); links to MDS working group
- Fergal Martin from Ensembl - ongoing plans for salmonid genomes (10-15 mins)

1300 - 1400: *Lunch*

1400 – 1515: Breakout groups

Questions and discussion points will include:

- **What do you want to get out of FAASG going forwards?**



- **Is funding for centralized coordination of FAASG essential?**
- **Should FAASG simply merge into FAANG?**
- **Is there interest in an update publication (FAASG whitepaper 2.0)?**
- **How to promote uptake of FAASG by industry?**
- **If FAASG had additional funding – how should it be spent?**

1515 - 1600: **Feedback from breakout** (10 mins per group); facilitators: TBC

1600 - 1700: **General discussion & wrap-up**; facilitators: TBC