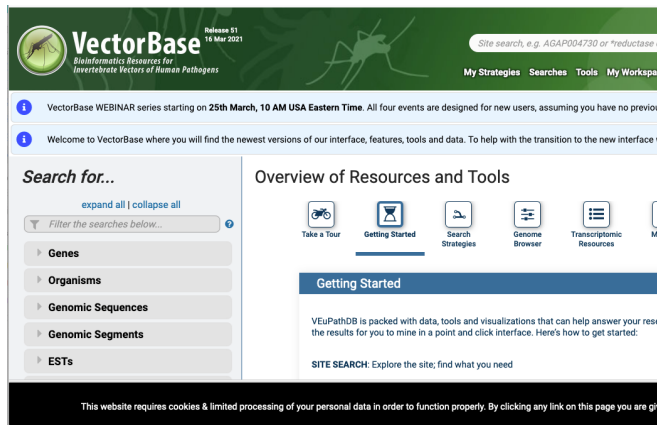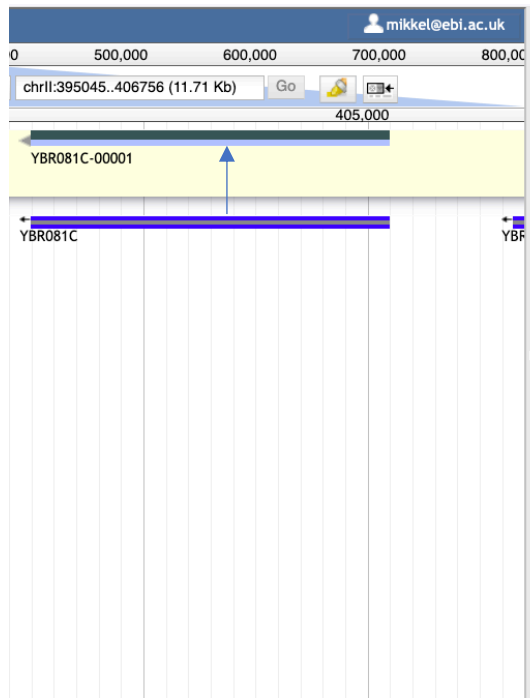1. Export existing Functional annotation from VEuParthDB in JSON.
2. Drag Gene to the annotation track as normal and select gene.
3. Import functional annotation via new Import function.
4. Select and Edit annotation as normal
5. Export as nornal