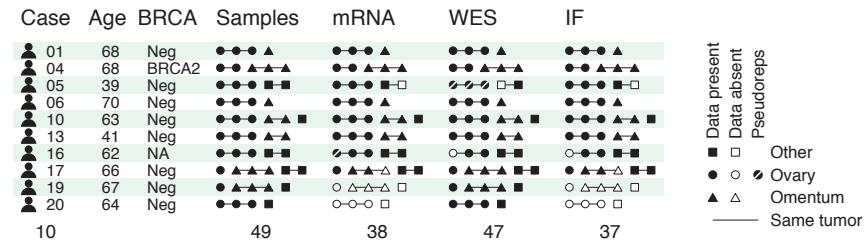
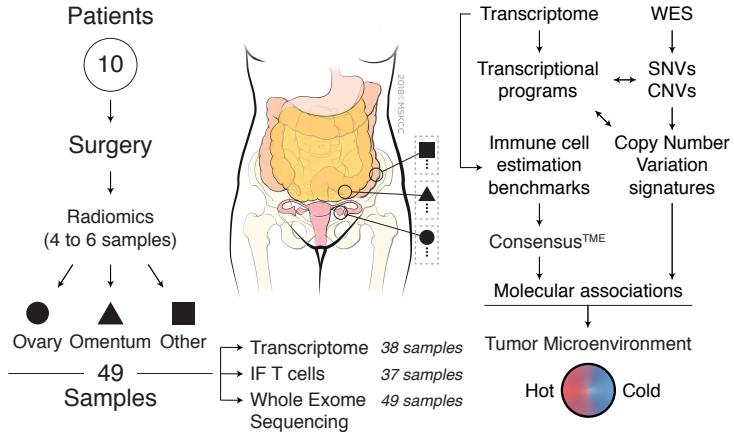
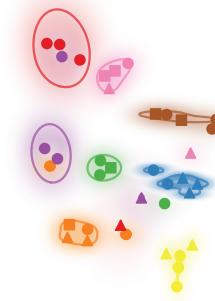


**a** Cohort I**b****c**

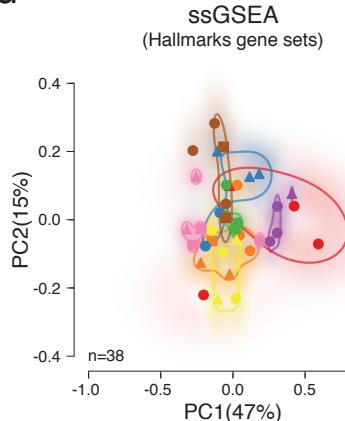
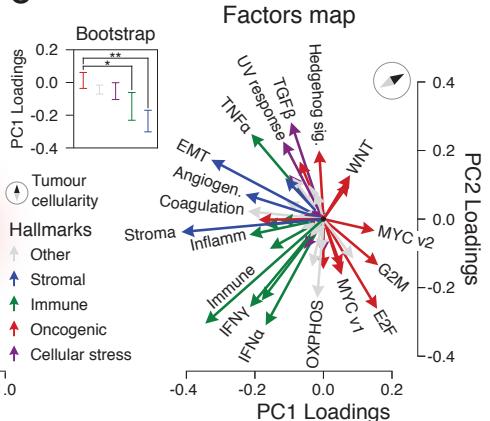
## t-SNE analysis of transcriptomes

n=38



Case

- 01
- 04
- 05
- 06
- 10
- 13
- 16
- 17

**d****e****f**