



Conquering Cancer Seminar:

Analysis of the initiation and progression of BRCA-mediated ovarian tumorigenesis at the single-cell level

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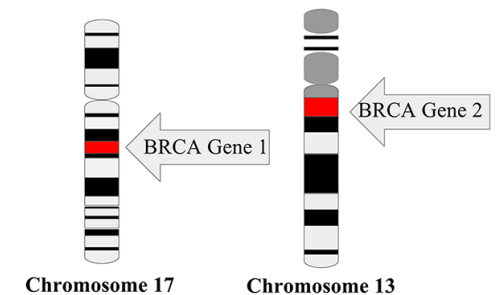
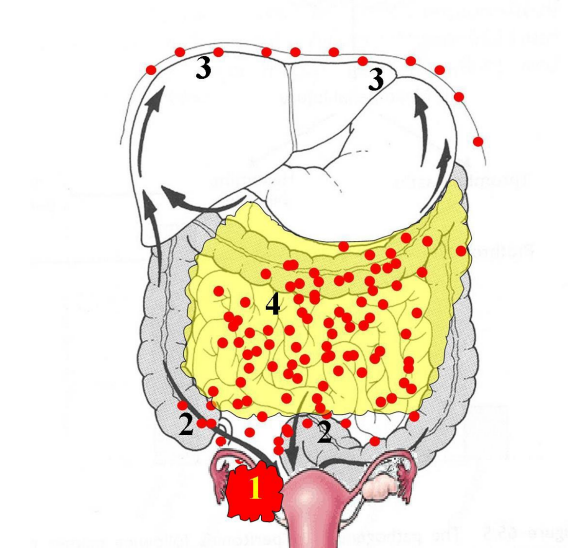
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Introduction

High grade serous ovarian cancer (HGSOC)

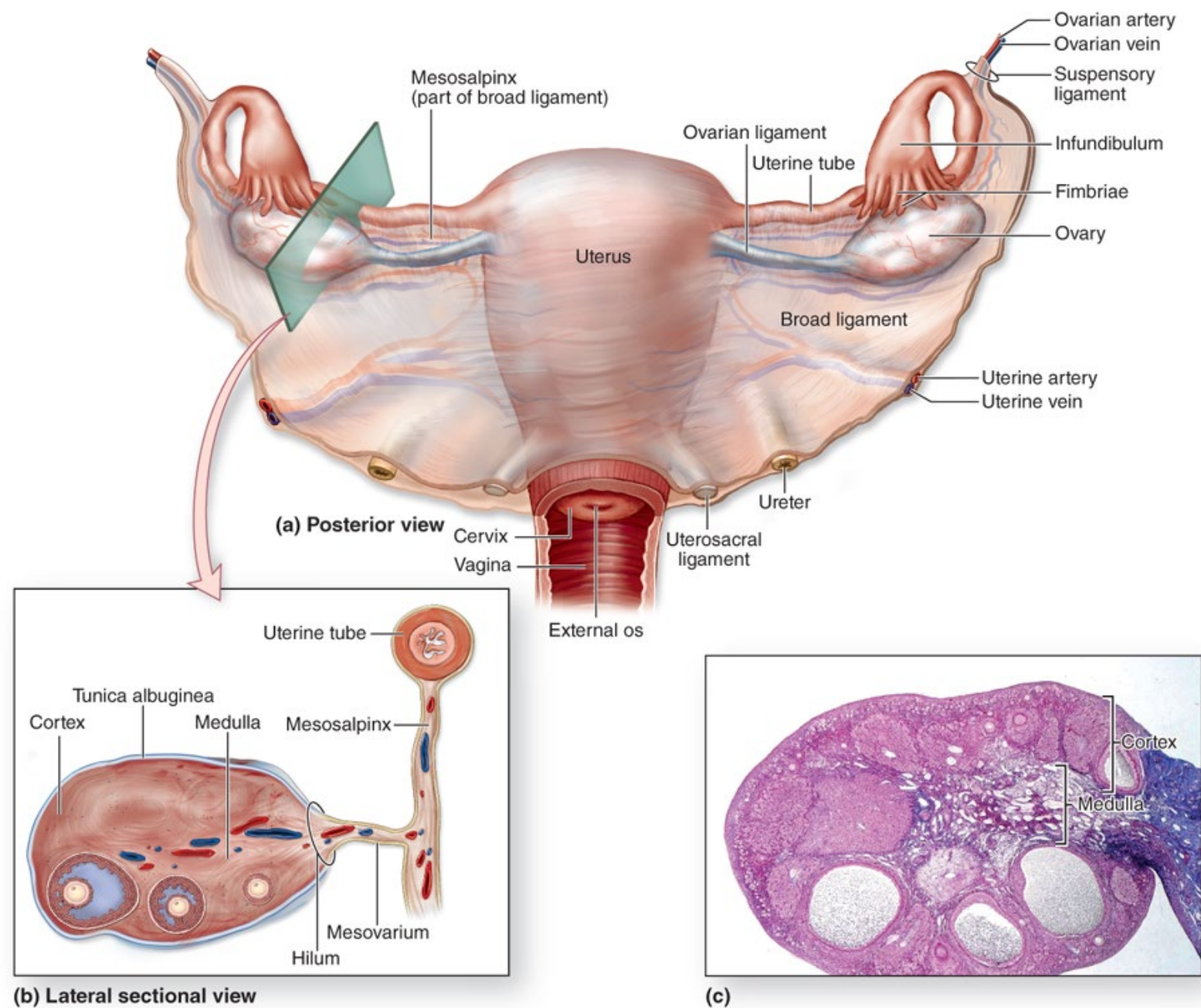
- HGSOC most common ovarian cancer, and most lethal Gyn cancer
 - **Presents at late stage and poor prognosis** – 5 yr survival ~ 35%; median survival ~3.5 years
 - **No screening/early detection**
 - **BRCA1/2**: 20-50% lifetime risk of epithelial ovarian cancer vs 1.8 % lifetime risk in the general population
 - 20% of HGSOC tumors have BRCA mutation
 - 13-15% = germline mut BRCA1/BRCA2, 7% somatic BRCA mut
- Knowledge gap
 - Lack knowledge about **disease initiation/early stages**
 - Current knowledge is based on whole tissue genomic data
 - Lack of information about cell-cell interaction/microenvironment



Background:

Ovarian Histology and Physiology

- **Cortex**
 - Surface epithelium
 - Oocytes
 - Stromal cells
- **Medulla**
 - Stromal cells, fibroblasts, vasculature
- **Cell types**
 - **Epithelial cells**
 - Endothelial, fibroblasts
 - Immune cells
 - Oocytes
 - Granulosa cells
 - Theca cells
- **Function:**
 - Harbors/ produces oocytes and sex hormones



OVARIAN EPITHELIAL CANCER

- High-Grade Serous Carcinomas
- Low-Grade Serous Carcinomas
- Clear cell carcinoma
- Endometrioid
- Mucinous

OVARIAN SEX CHORD-STROMAL TUMORS

Stromal tumors

1. Fibroma
2. Thecoma
3. Fibrosarcoma
4. Leydig cell tumor
5. Steroid cell tumor
6. Sclerosing stromal tumor

Sex chord tumors

1. Adult granulosa cell tumor
2. Juvenile granulosa tumor
3. Sertoli cell tumor
4. Sex chord tumor with annular tubules

Mixed sex chord-stromal tumors

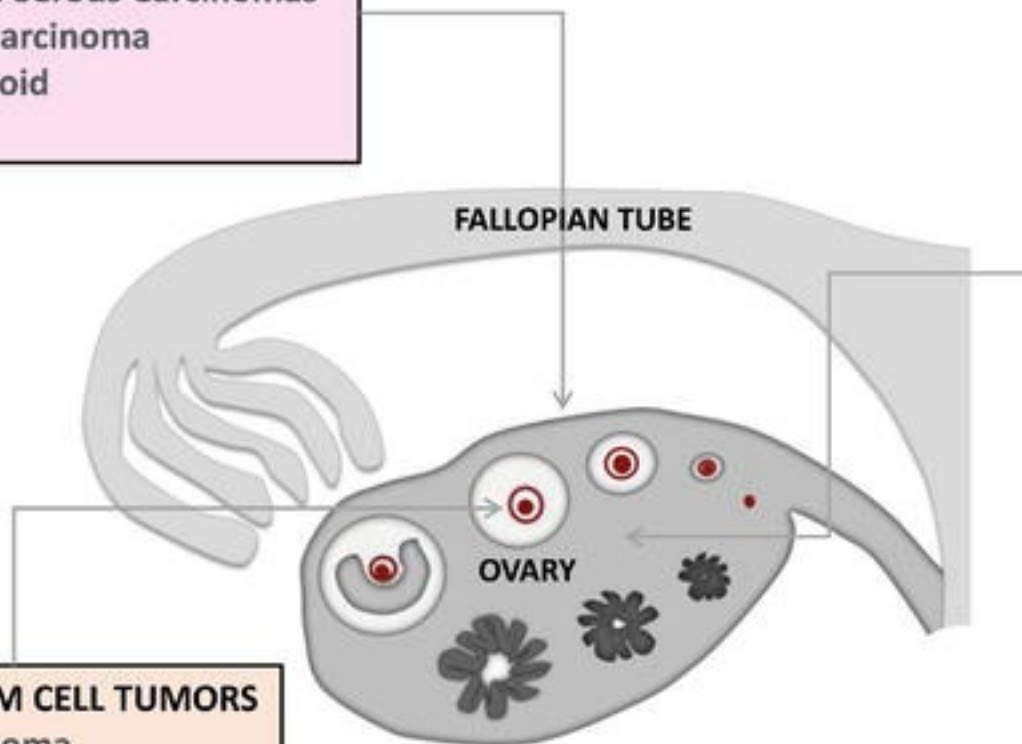
1. Sertoli-Leydig cell tumor

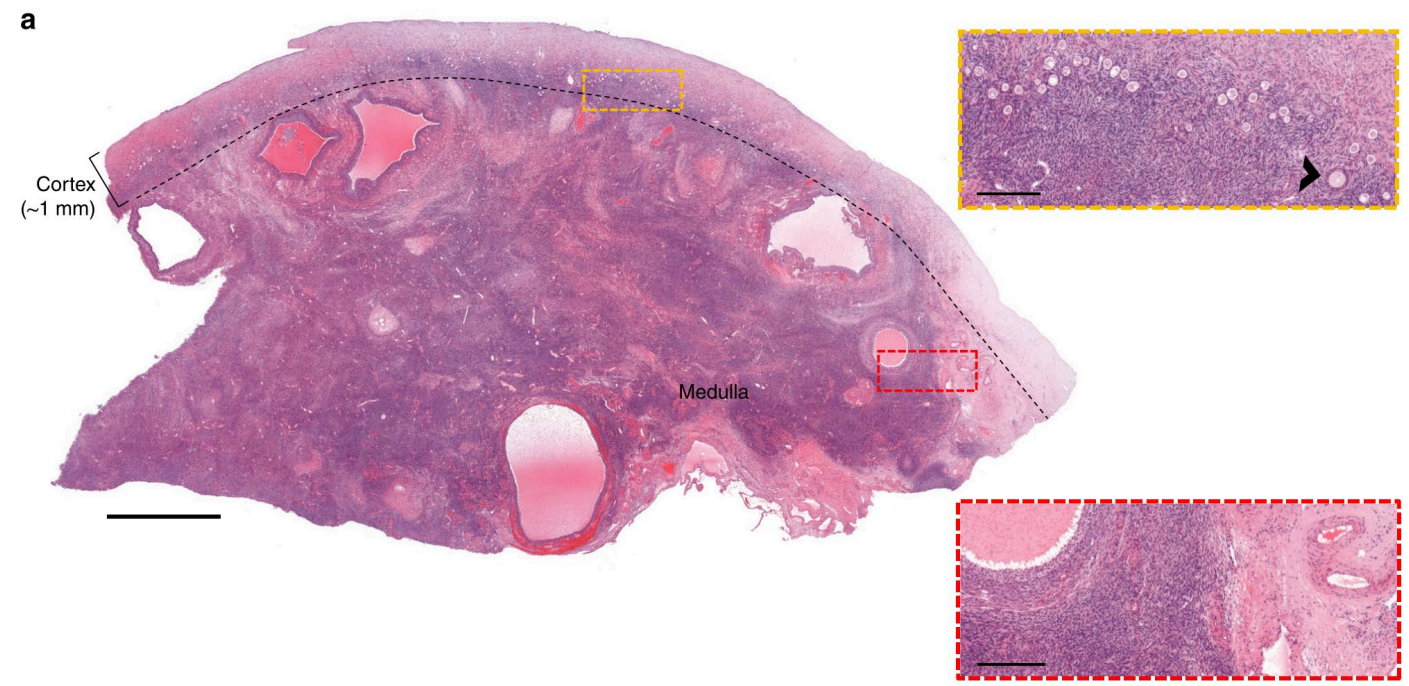
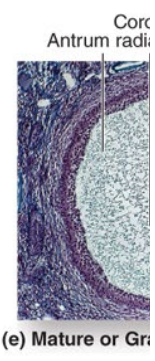
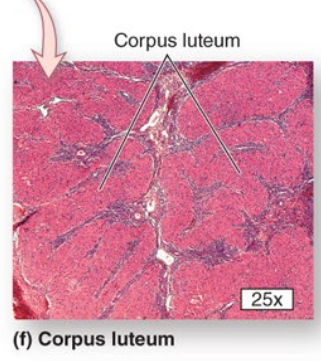
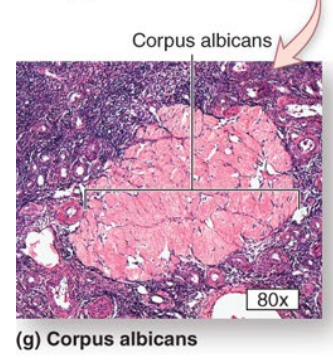
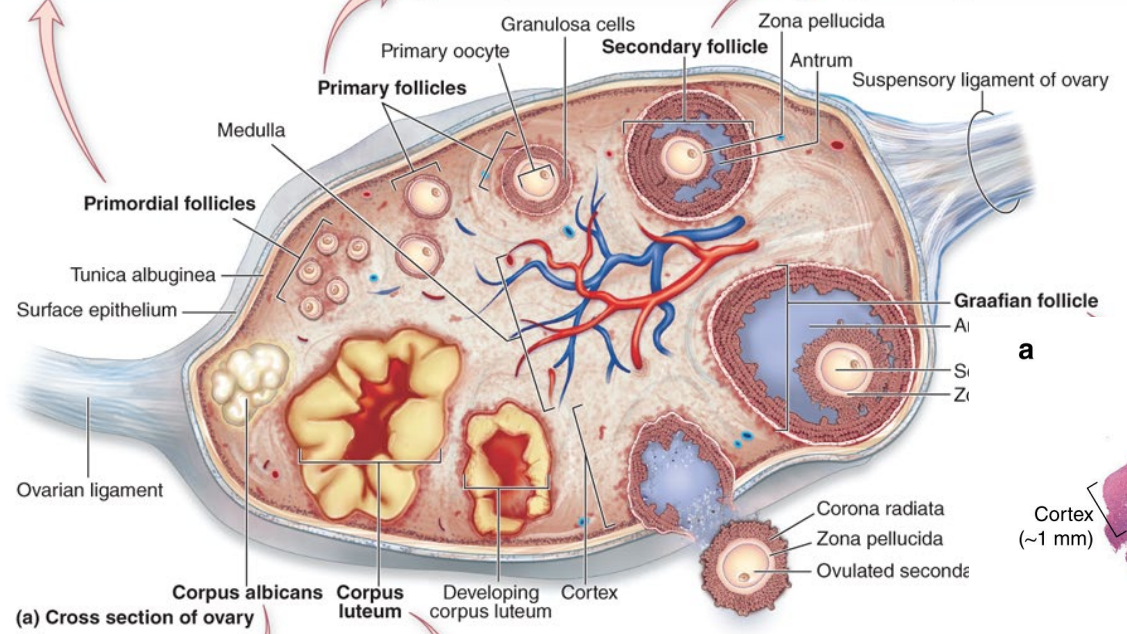
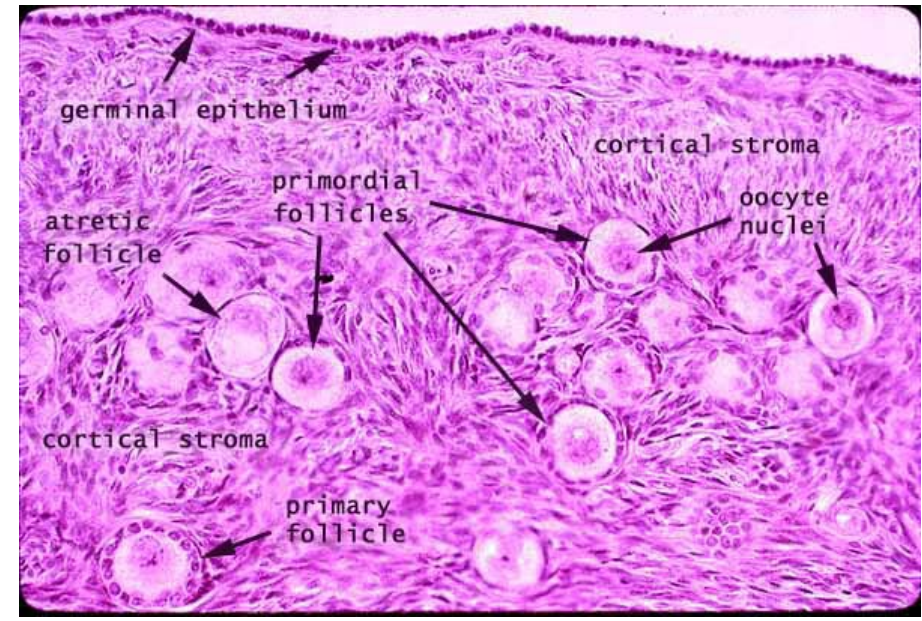
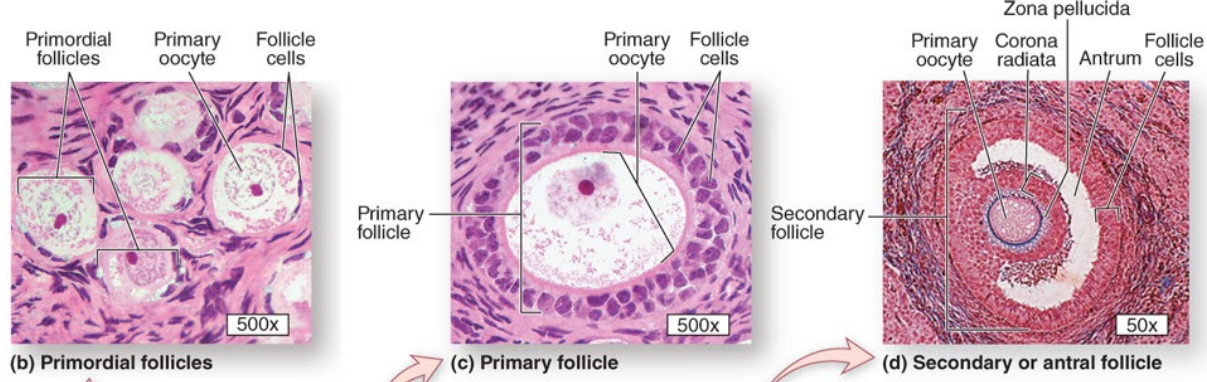
OVARIAN GERM CELL TUMORS

1. Dysgerminoma
2. Immature teratoma
3. Yolk sac tumors
4. Mixed germ cell tumors

SMALL CELL CARCINOMA OF THE OVARY

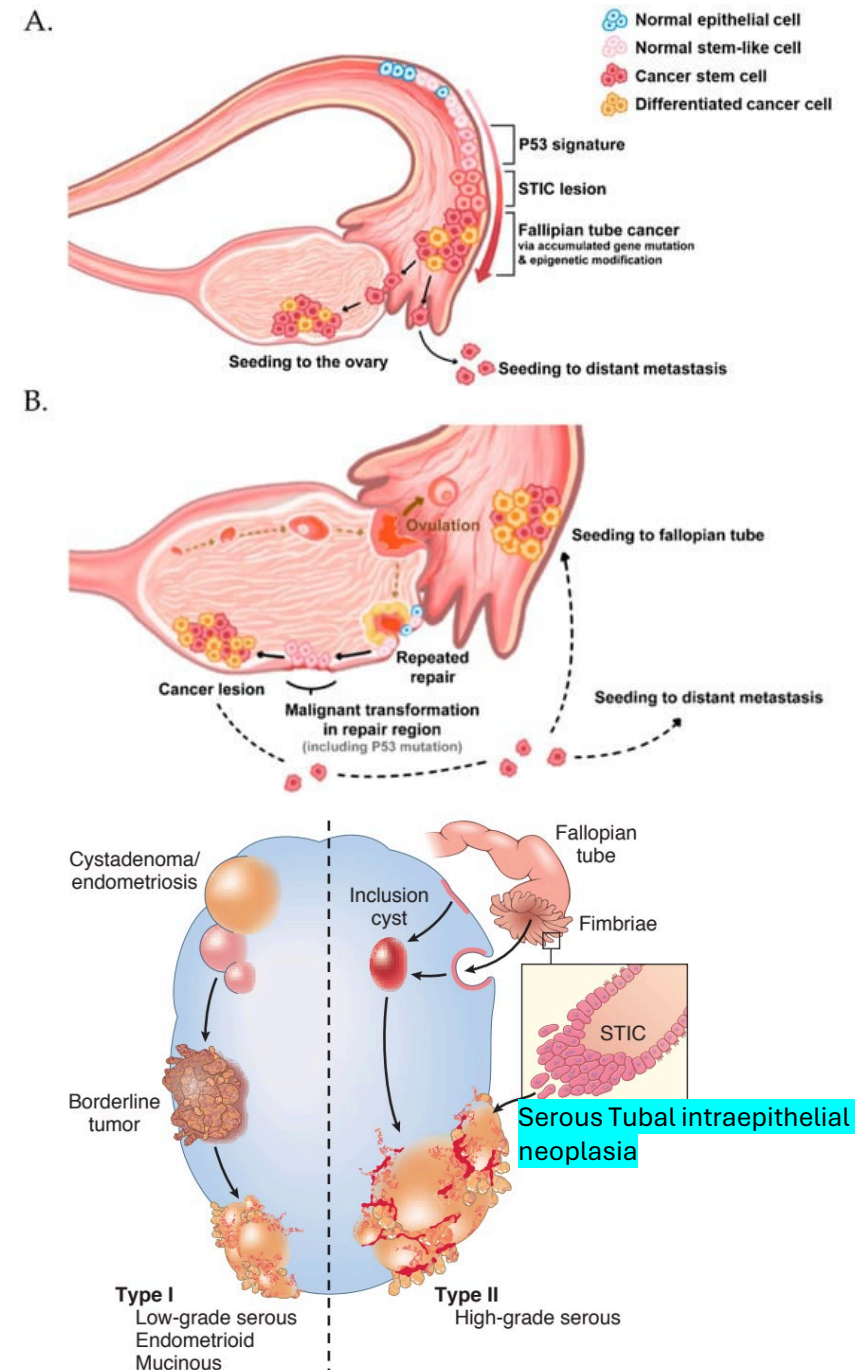
1. SCCO - hypercalcemic type
2. SCCO - pulmonary type





Background: Pathophysiology

- Development of ovarian cancer
 - **Genomic instability**, TP53
 - **Incessant ovulation**:
 - Recurrent ovulation as a transforming event → damage to epithelium
 - **Primary site of origin? + inclusion cyst**:
 - extra-ovarian epithelia (endometrium or fallopian tube epithelium)
 - endometriosis or fallopian tube epithelium seeded on the ovary as inclusion cysts
 - transformed epithelial cells seed the ovary and its microenvironment → neoplasia



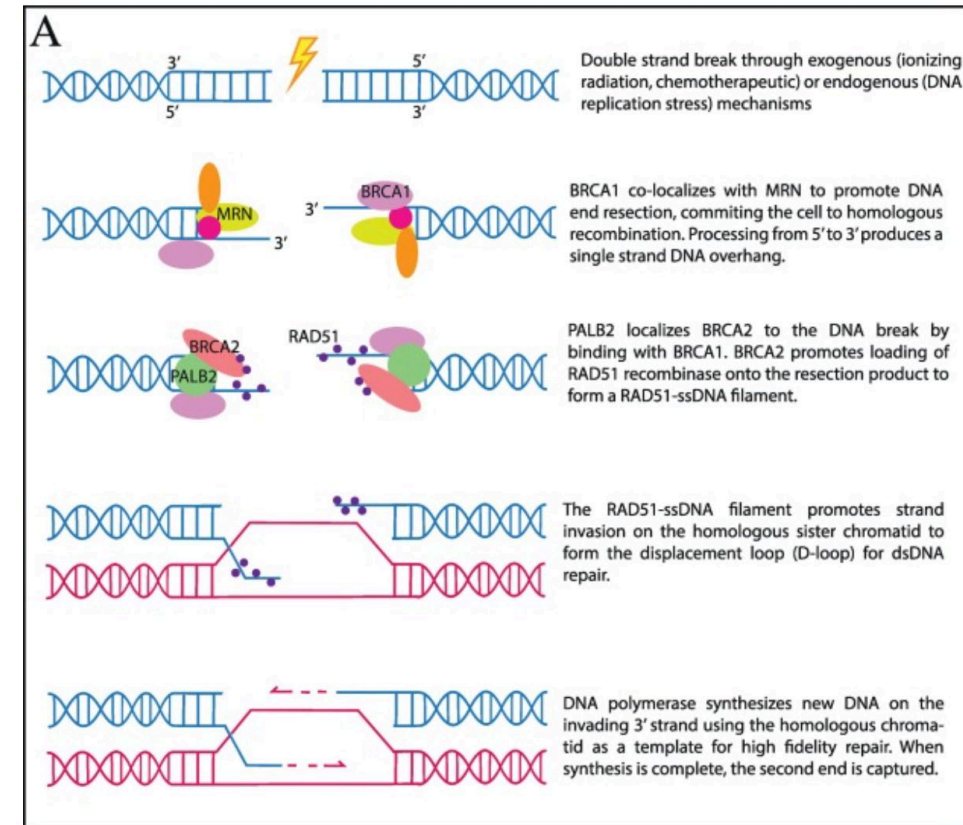
Genetic susceptibility: BRCA mutation / HRD

= ovarian cancer susceptibility genes

BRCA1 = 50% OC risk

BRCA2 = 20% OC risk

- **BRCA1** and **BRCA2** = proteins involved in **homologous recombination DNA repair**
 - Homologous recombination (HR) is one method to repair double-strand DNA breaks (DSB)
- 50% of tumors possess mutations in HR genes = homologous recombination deficiency or HRD, which includes the BRCA genes



Background:

Treatment and clinical course/prognosis

- **Treatment**

- Cytoreductive surgery
- Carboplatin/Paclitaxel +/- Bevacizumab – 6 cycles, ? Maintenance therapy

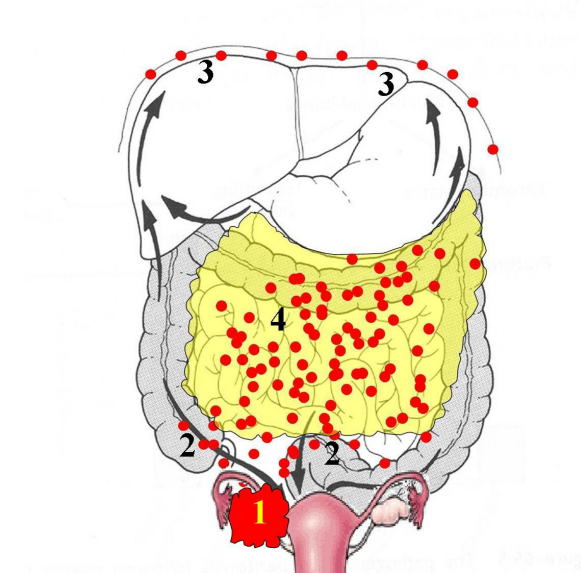
- **Clinical course/Prognosis**

- 75% present at stage 3+
- 65% recurrence within 2 years, 75% at 3 years
- 5-year cause-specific survival: stage 3 (42%) stage 4 (26%)
- Fast growth of tumor nodules encasing organs → ascites, carcinomatosis, bowel obstruction

- **Improvements**

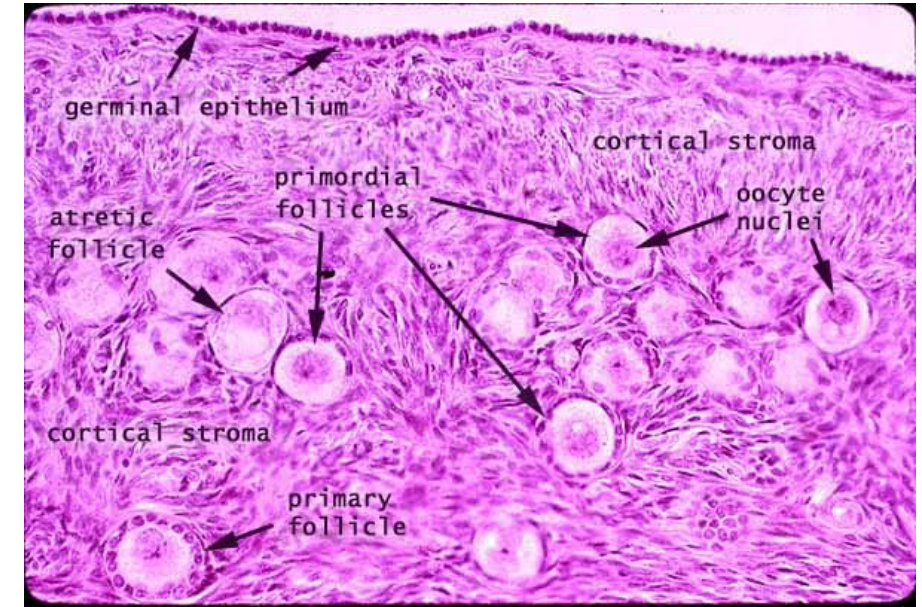
- 5-10% improvement in survival rate over 30 years
- Maintenance therapies

- **Still NO screening or early detection**

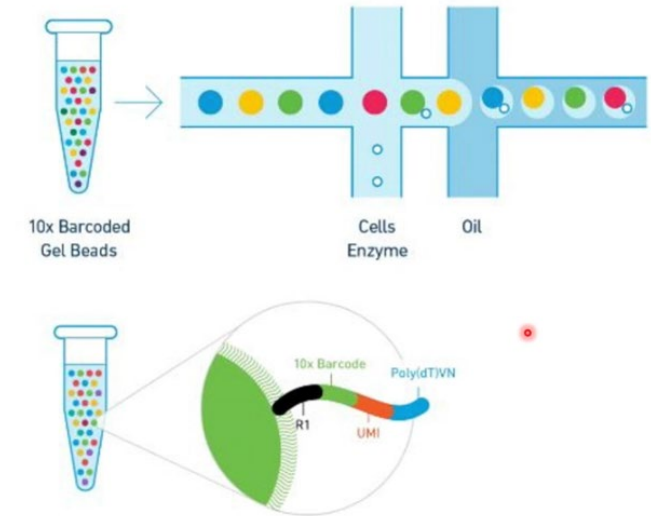


Aim / hypothesis

- Identify cell populations, RNA expression patterns and cell/cell interactions within the ovary that lead to changes in epithelial cells that can cause ovarian cancer
- We hypothesize that BRCA+/mut carriers exhibit alterations in ovarian epithelial and stromal cell populations including changes in genetic expression, concentrations, and cellular interactions, that promote epithelial cell changes leading to cancer initiation.



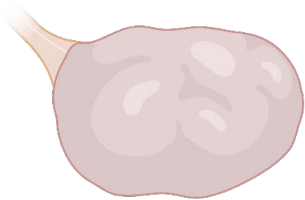
- Translational discovery:
 - These cell populations and RNA expression pattern changes could identify potential:
 - Precursor lesions/cell types
 - Biomarkers for pre-cancerous lesions and prevention techniques
 - Therapeutic targets



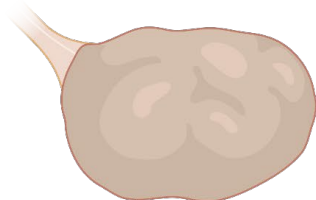
Methods

- **Tissue procurement**
 - FFPE archived oophorectomy tissue
 - **Benign oophorectomy BRCA wildtype**
 - **Pre-neoplastic/prophylactic BRCA mut**
 - **Ovarian Tumor BRCA mut**
 - **Met Tumor BRCA mut**
- **Single-cell RNA sequencing**
- **Spatial omics and multiplexed imaging**

Normal Ovary
(control)



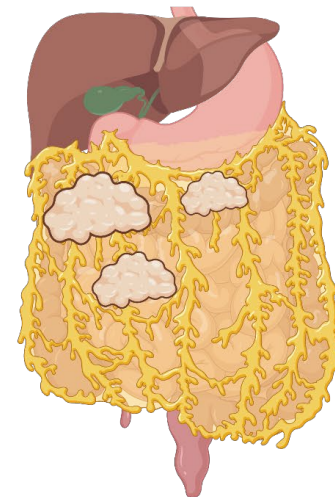
Pre-neoplastic
BRCA+/mut ovary



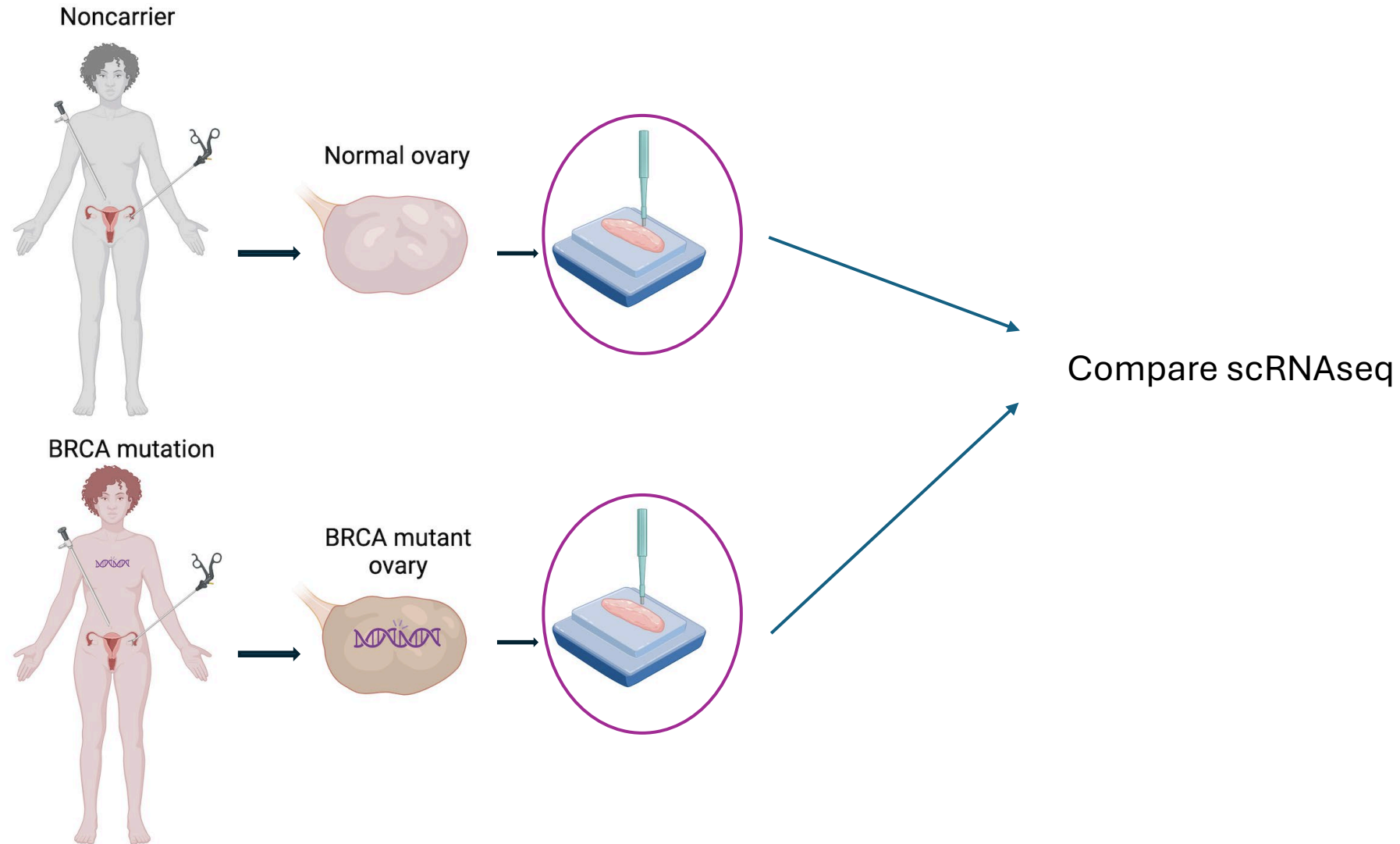
Ovarian BRCA+/
mut tumor



Metastatic BRCA
+/mut tumor

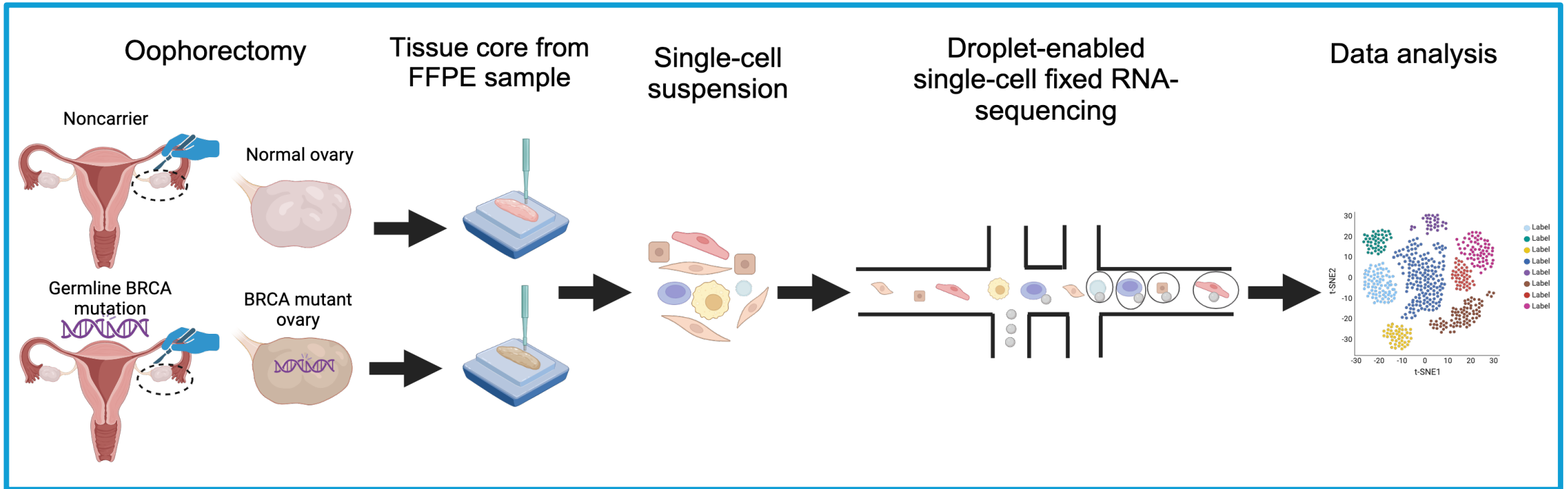


Normal ovary vs BRCA carrier ovary



Single cell RNA sequencing

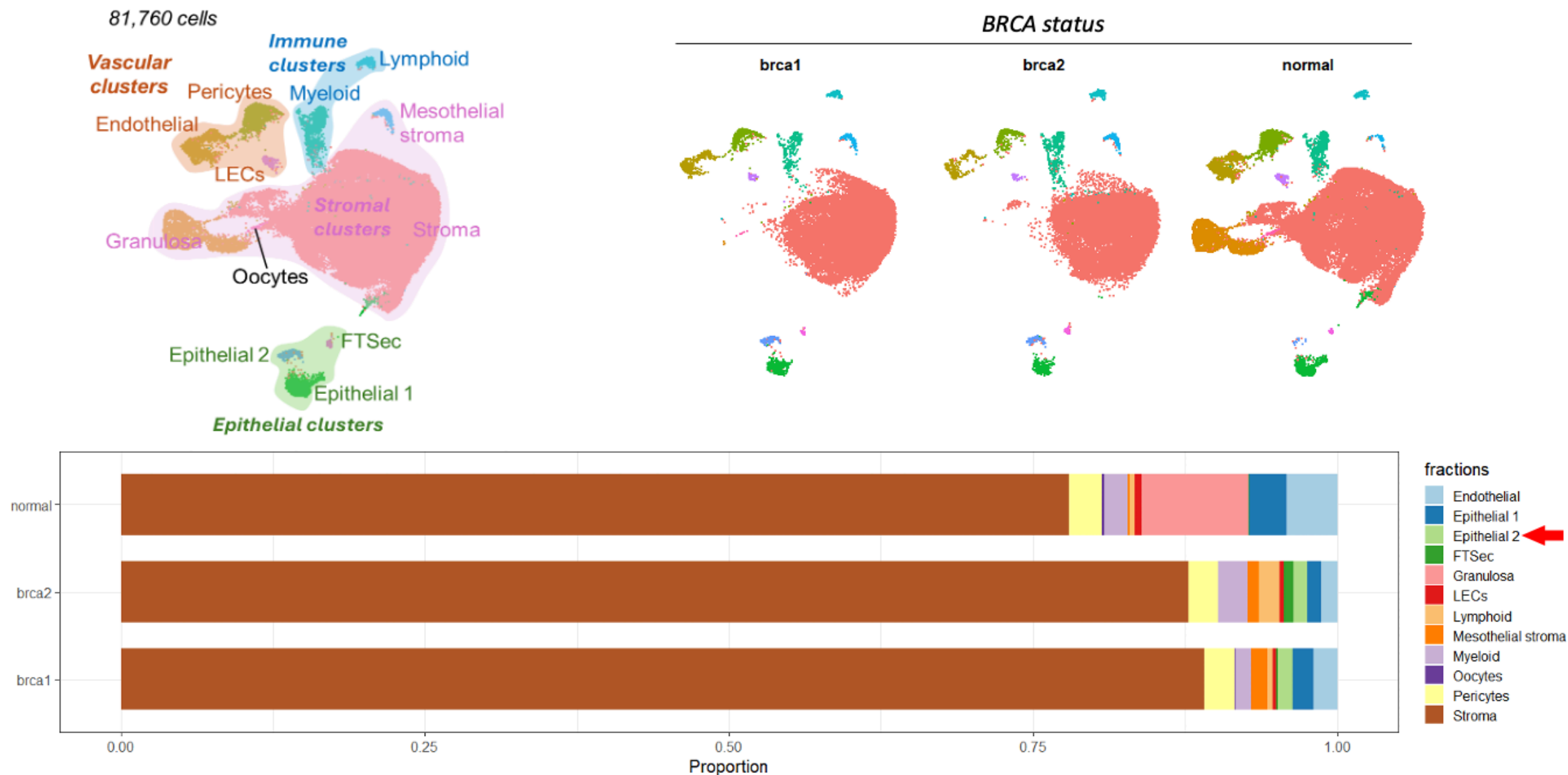
Single cell RNA seq workflow



Genotype	No. of samples
Noncarrier	8
BRCA1 mut	5
BRCA2 mut	5

Data analysis

Figure 2. Cell cluster types and frequencies in Normal vs BRCA+/mut ovarian



Marker genes for epithelial 2

