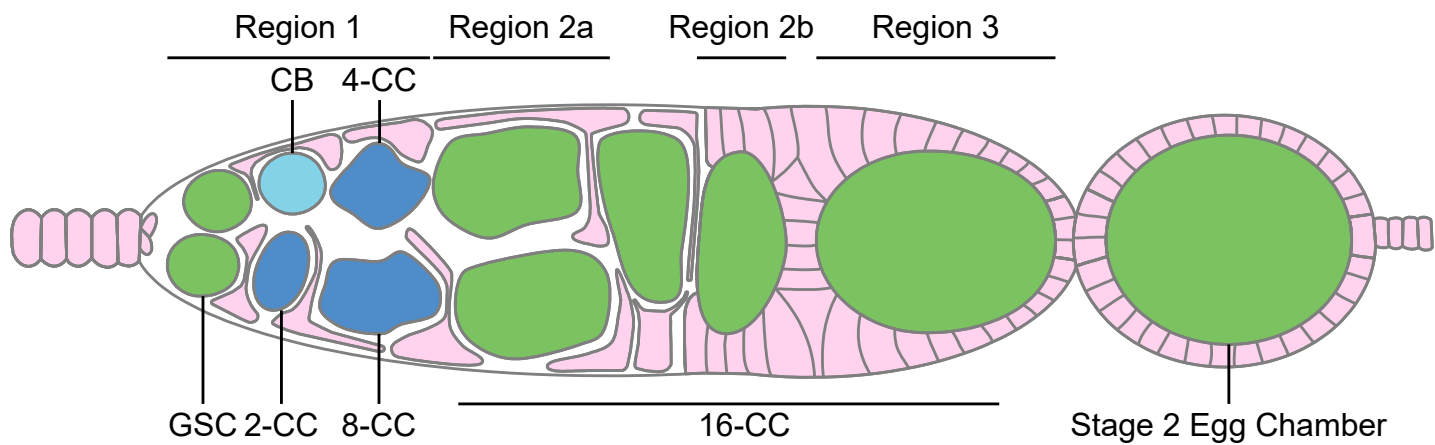


A



B

Screenshot of the Oo-site web application interface. The interface is divided into a left sidebar and a main content area. The sidebar contains the following elements:

- Display Cartoon** (with a red circle 2 next to it)
- Select a dataset** (with a red circle 3 next to it)
- View dataset:**
 - ☒ Input mRNAseq
 - ☐ Polysome-seq
 - ☐ Single Cell-seq: Germline
 - ☐ Single Cell-seq: Soma
- Developmental Progression** (with a red circle 3 next to it)
- Heatmap**
- Gene Groups**
- Generate report** (with a red circle 5 next to it)
- Take a tour!** (with a red circle 1 next to it)

The main content area displays a cartoon of the embryo with a color-coded expression scale (Binned Expression) ranging from None to Very High. The scale is color-coded: None (grey), Very Low (light blue), Low (blue), Med (dark blue), High (cyan), and Very High (magenta). The cartoon shows the expression of the gene of interest (RpS19b) across different developmental stages and regions. The main content area also includes a **Controls** panel (with a red circle 4 next to it) containing the following options:

- Data set:** Symbol
- Gene of Interest:** RpS19b
- ☒ Display Expression Values
- ☒ Display Stage Labels