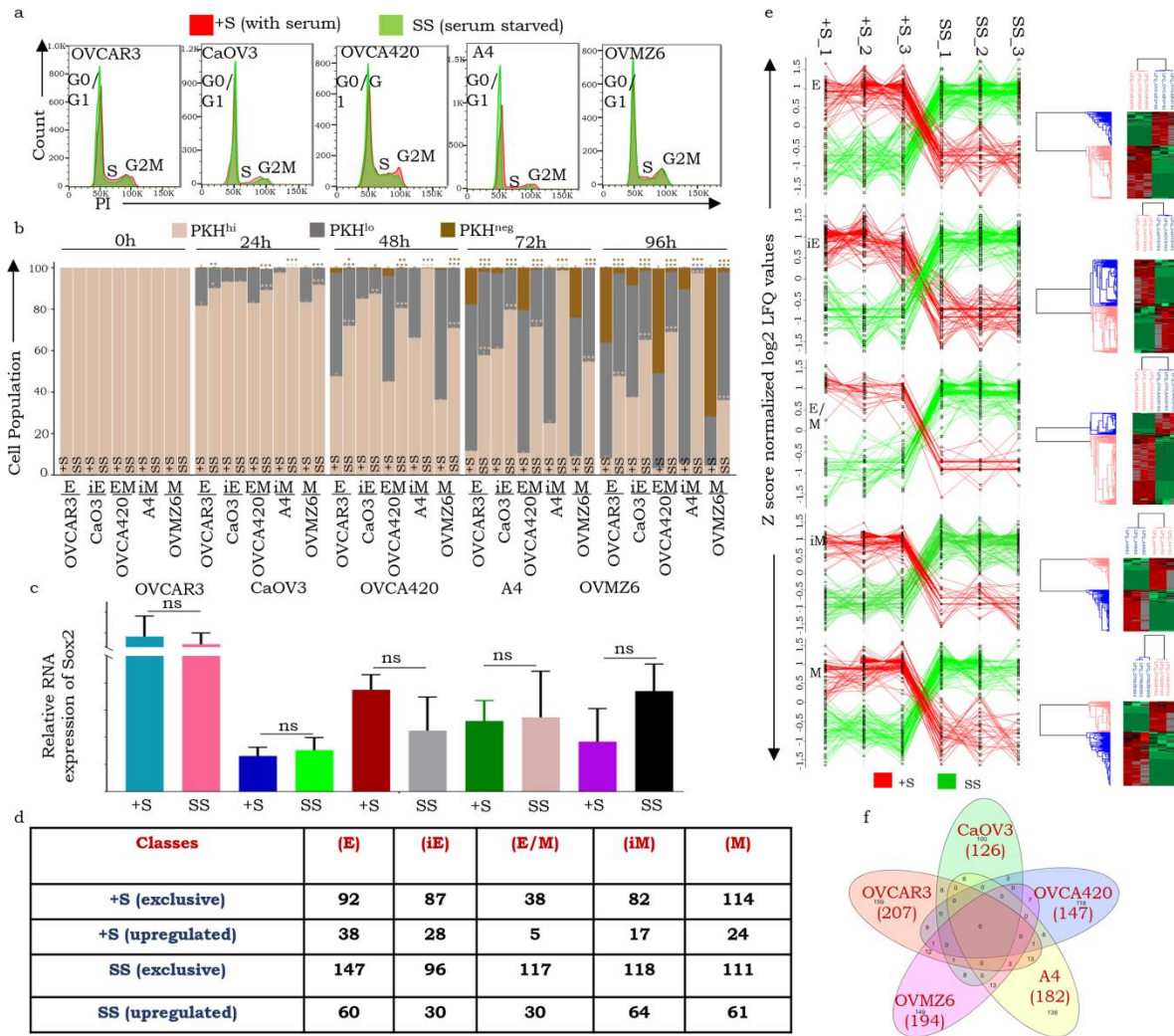


Supplementary Fig. 1

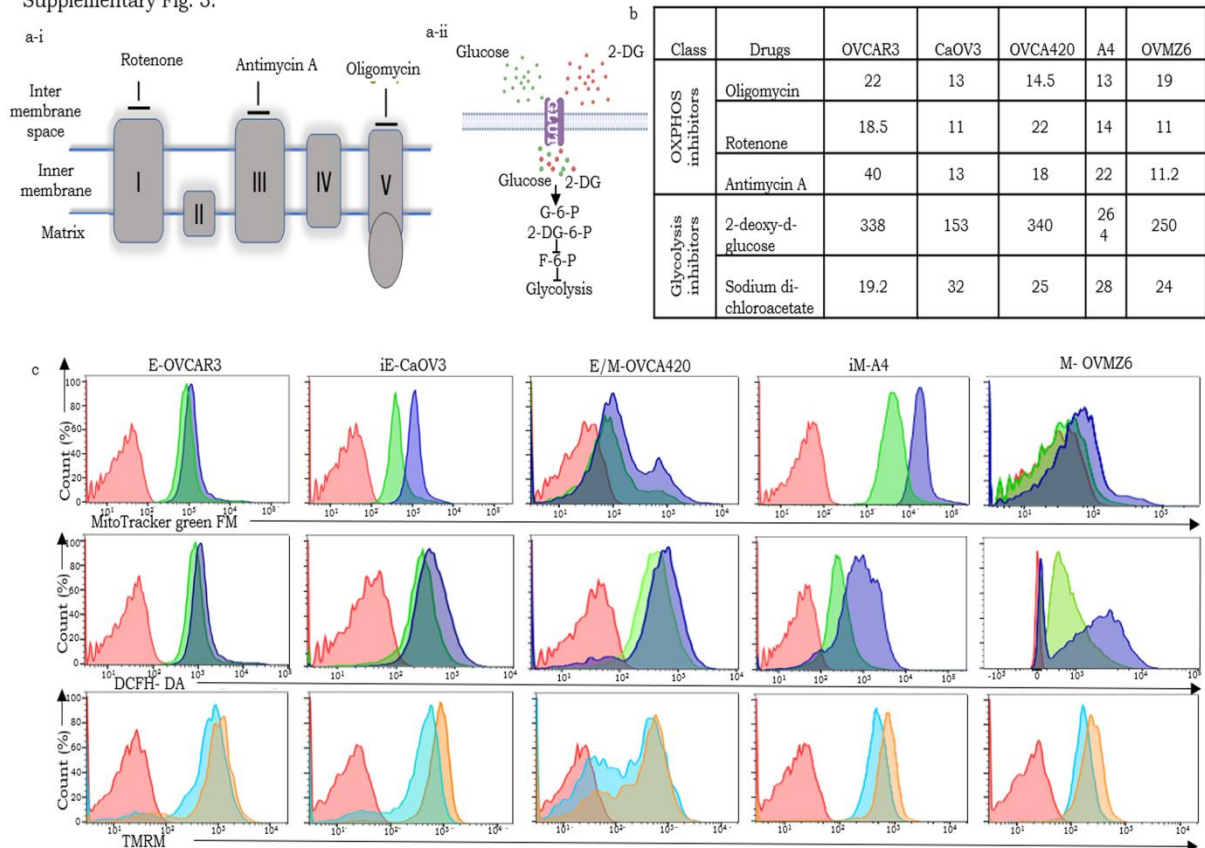


Supplementary Fig.1. a. Representative histogram indicating cell cycling of HGSC cells grown under SS (96h) and +S conditions (green-SS, red-+S); **b.** Stacked-bar graph indicating the PKH^{hi} PKH^{lo} and PKH^{neg} fractions in HGSC +S and SS derivatives at different time interval (0h, 24h, 48h, 72h and 96h); **c.** Bar graph indicating relative levels of Sox2 expression in HGSC phenotypes under +S and SS conditions; **d.** Table consolidating the number of upregulated and exclusive proteins under SS and +S conditions across the phenotypic spectrum (E- Epithelial-OVCA420, iE-intermediate Epithelial-CAOV3, E/M-Epithelial/Mesenchymal hybrid-OVCA420, iM- intermediate Mesenchymal-A4, M-Mesenchymal-OVMZ6); **e.** Left panel- profile plot representing SS and +S enriched proteins, right panel- heat map of the same; **f.** Venn-diagram indicating shared and exclusive proteins between HGSC phenotypes under conditions of serum starvation (SS). **p < 0.05, ***p < 0.01 and ****p < 0.001.

[illegible]

Supplementary Fig. 2. Representative cluster plots and pie diagram indicating enriched GO biological process and cellular components in HSGC phenotypes following SS conditions. Regardless of the cellular phenotype, serum starvation response engages different mitochondrial-related proteins

Supplementary Fig. 3.



Supplementary Fig. 3. HGSC cells prefer OXPHOS under SS accompanied by significantly increased active mitochondria. **a-i, a-ii.** Schematic indicating site of action of OXPHOS and glycolysis inhibitors respectively; **b.** IC₅₀ values of the specific inhibitors on HGSC cell lines; **c.** Histograms representing, mitochondrial mass (MitoTracker green FM-upper panel), ROS (DCFH-DA-middle panel) and mitochondrial potential/ activity (TMRM- lower panel) at 72h under SS & +S conditions. Violet and green colour in the upper and middle panels represent SS and +S histogram respectively, while the same is represented by orange and blue, respectively, in the lower TMRM panel.

The diagram illustrates the effect of Etomoxir on mitochondrial fatty acid oxidation (FAO) and the resulting metabolic state in two conditions: +S (Saturated) and SS (Starvation).

+S Condition (Left):

- The cell contains a large nucleus and a large mitochondrion.
- FAO is active, converting Acyl CoA to Acetyl CoA, which enters the TCA cycle, producing NADH/FADH₂ and driving the ETC.
- Etomoxir inhibits CPT1, which is responsible for the transport of Acyl CoA into the mitochondrion.
- The cell is labeled "Less sensitive to Etomoxir".

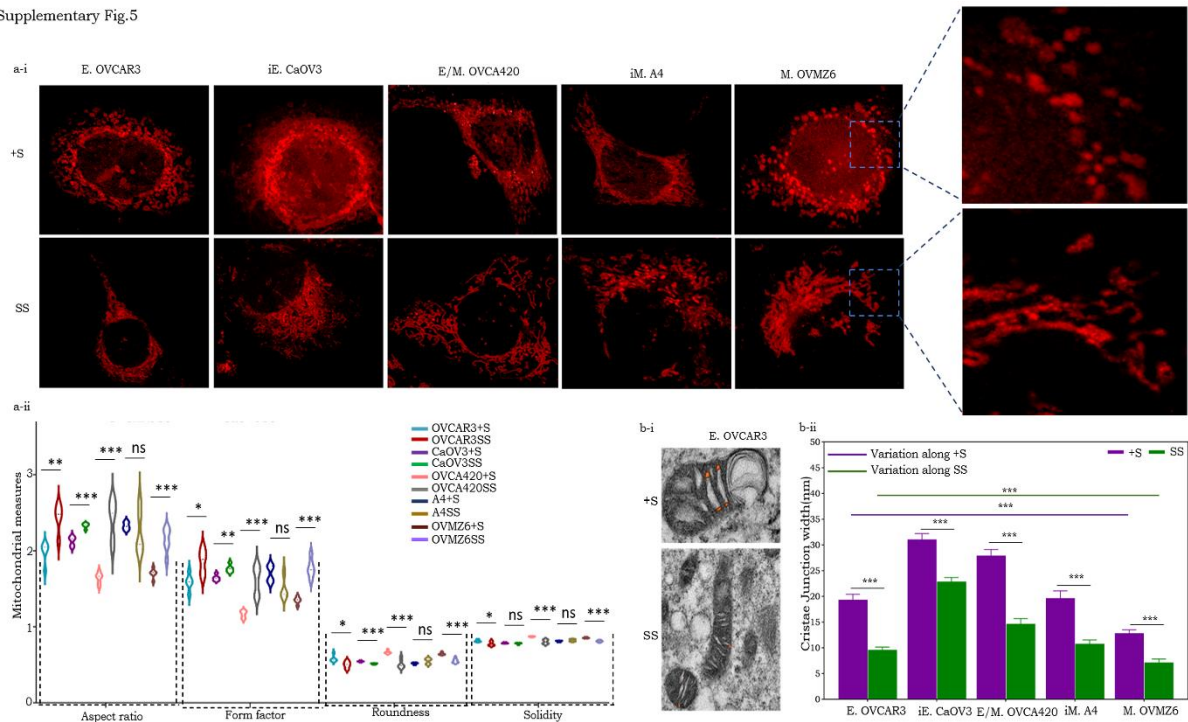
SS Condition (Right):

- The cell contains a large nucleus and a smaller mitochondrion.
- FAO is inhibited, leading to a decrease in Acetyl CoA, TCA cycle activity, and ETC activity.
- Etomoxir inhibits CPT1, which is responsible for the transport of Acyl CoA into the mitochondrion.
- The cell is labeled "More sensitive to Etomoxir".

Summary of Effects:

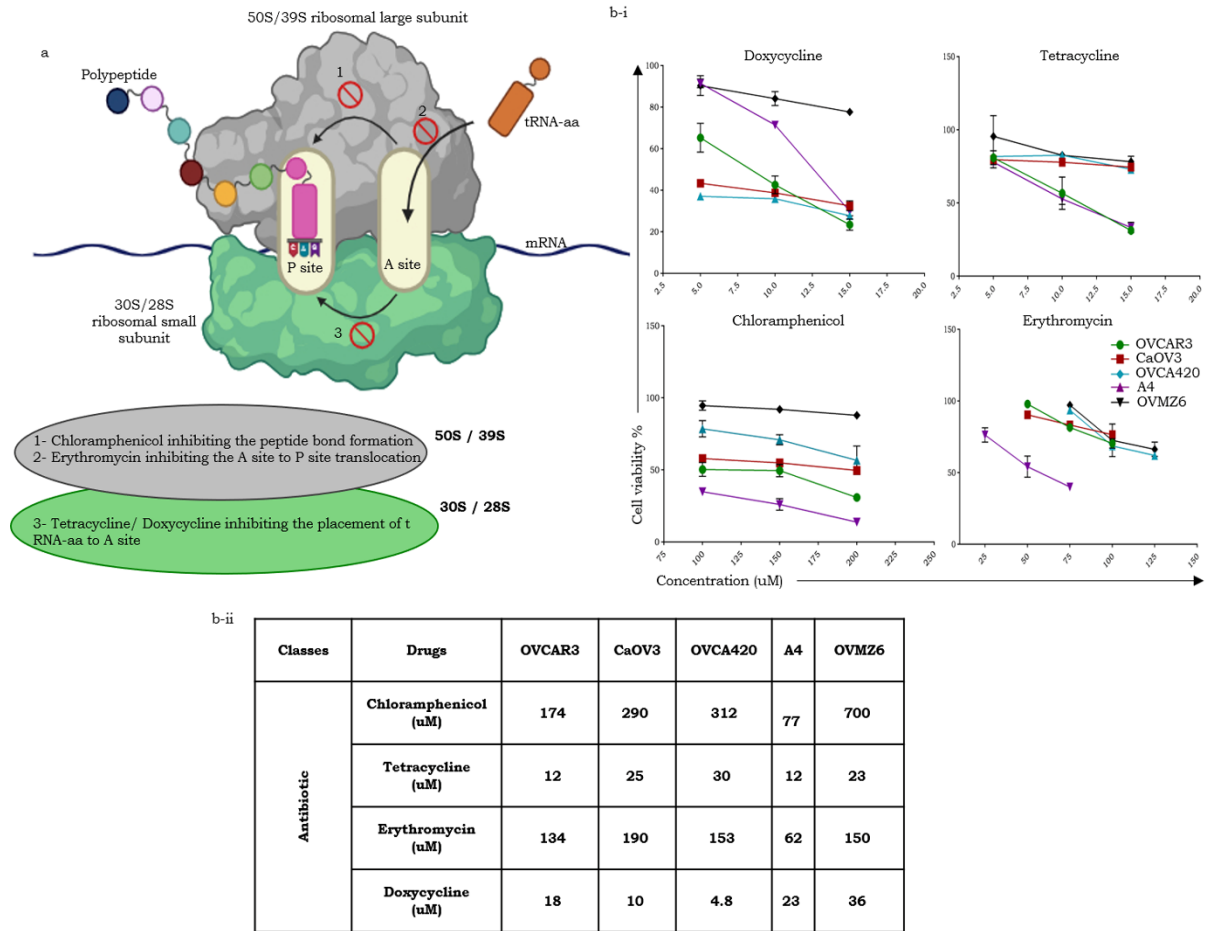
Parameter	+S Condition	SS Condition
Lipolysis	↓	↑
No. of Lipid droplets	↑	↓
Sensitivity to Etomoxir	↓	↑
FAO	↓	↑
TCA	↓	↑

Supplementary Fig.5



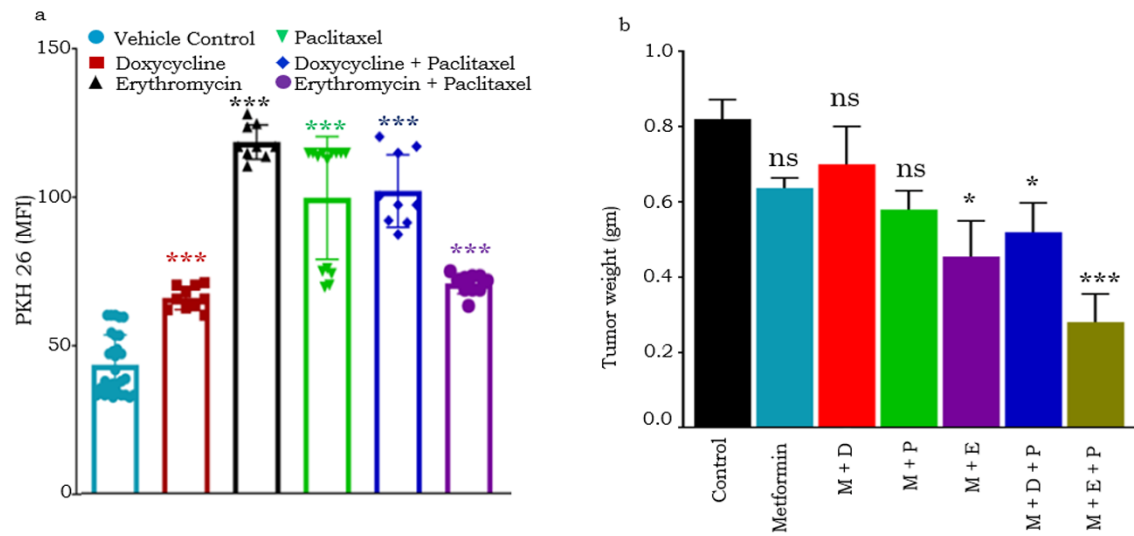
Supplementary Fig. 5. Under SS conditions, HGSC cells mitochondria undergo fusion to fulfil the energy demands necessary for their survival: **a-i**. MitoTracker deep red confocal microscopy images indicating the mitochondrial morphology across HGSC phenotypes in the presence (upper panel) and absence (lower panel) of serum; **a-ii**. Violin plot representing the mitochondrial structural parameters analysed on ImageJ on 48h SS vs +S samples (derived from at least 50 individual cells per condition across the various phenotypes); **b-i**. Representative confocal images highlighting CJs (orange in colour) in 48h +S and SS OVCAR3 cells; **b-ii**. Bar-graph indicating CJ width in HGSC cells under SS and +S conditions. **p < 0.05, *p < 0.01 and ***p < 0.001.

Supplementary Fig.6



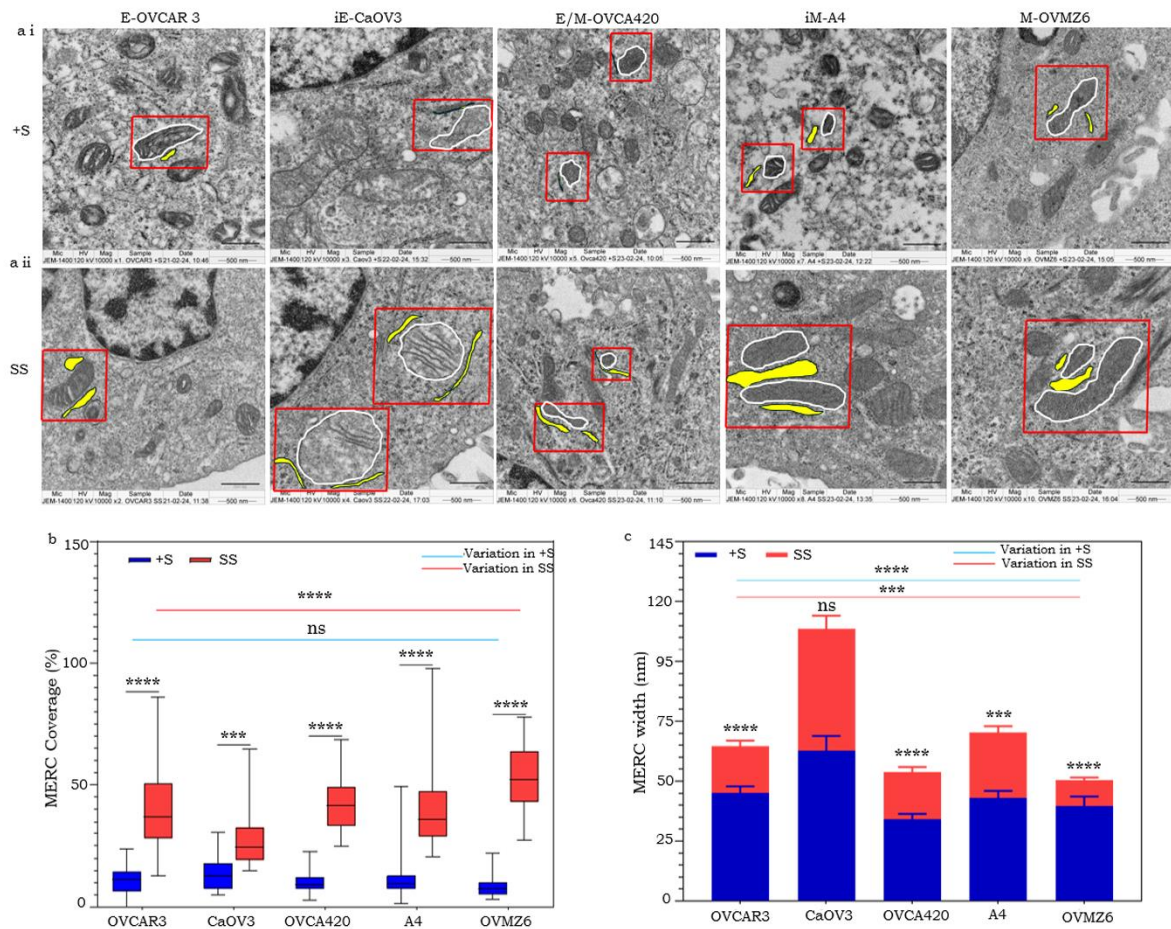
Supplementary Fig. 6. Site and mechanism of action of antibiotics and their IC₅₀ values in different HGSC phenotypes. **a.** Schematic representation of site and mode of action of different classes of antibiotics; **b-i.** Line plots representing IC₅₀ value determination of antibiotics in different HGSC phenotypes; **b-ii.** Table indicating the computed IC₅₀ values of antibiotics in different HGSC cell lines.

Supplementary Fig.7.



Supplementary Fig. 7. Tumor inhibitory effects of OXPHOS and mitochondrial translation inhibitors. **a.** Bar graph indicating PKH quenching in A4 tumors harvested at Day 21 of different drug regimens; **b.** Bar-graph indicating A4 tumor weights at Day 21 of different drug regimens. **p < 0.05, *p < 0.01 and ***p < 0.001.

Supplementary Fig.8



Supplementary Fig. 8. Serum starvation led to reduced mitochondrial-endoplasmic-reticulum contact (MERC) distance across all phenotypes. **a.** Representative 2D-TEM images highlighting the MERC distance under +S (upper panel) and SS (lower panel) conditions; **b.** Floating bar-graphs indicating the MERC coverage in HGSC phenotypes under +S and SS conditions; **c.** Stacked bar-graph representing the MERC width across the phenotypes under +S (blue) and SS (red) conditions. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ and **** $p < 0.0001$

