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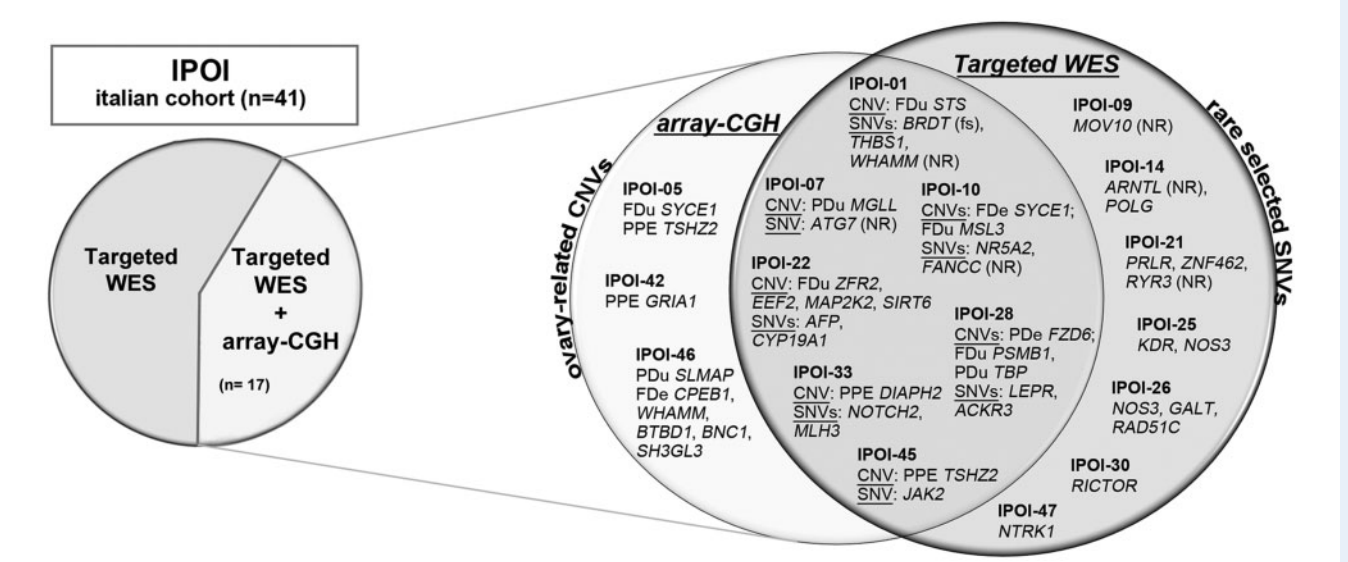
SUPPLEMENTARY DATA



Supplementary Figure S1. Results showing the integrated approaches (array-CGH and targeted-WES) performed on the Italian POI cohort. Among the 41 patients of the cohort, 17 were subjected to the combined method (Bestetti *et al.*, 2019), and the results are shown in the enlarged Venn diagram. For each patient, the obtained data are shown, and the intersection shows the patients positive for both rare ovary-related CNVs and SNVs/Indels. CGH, comparative genomic hybridisation; CNV, copy number variant; FDe, full deletion; FDu, full duplication; fs, frameshift; NR, never reported variant; PDe, partial deletion; PDu, partial duplication; POI, primary ovarian insufficiency; PPE, potential position effect; SNVs, single-nucleotide variants; WES, whole exome sequencing.