

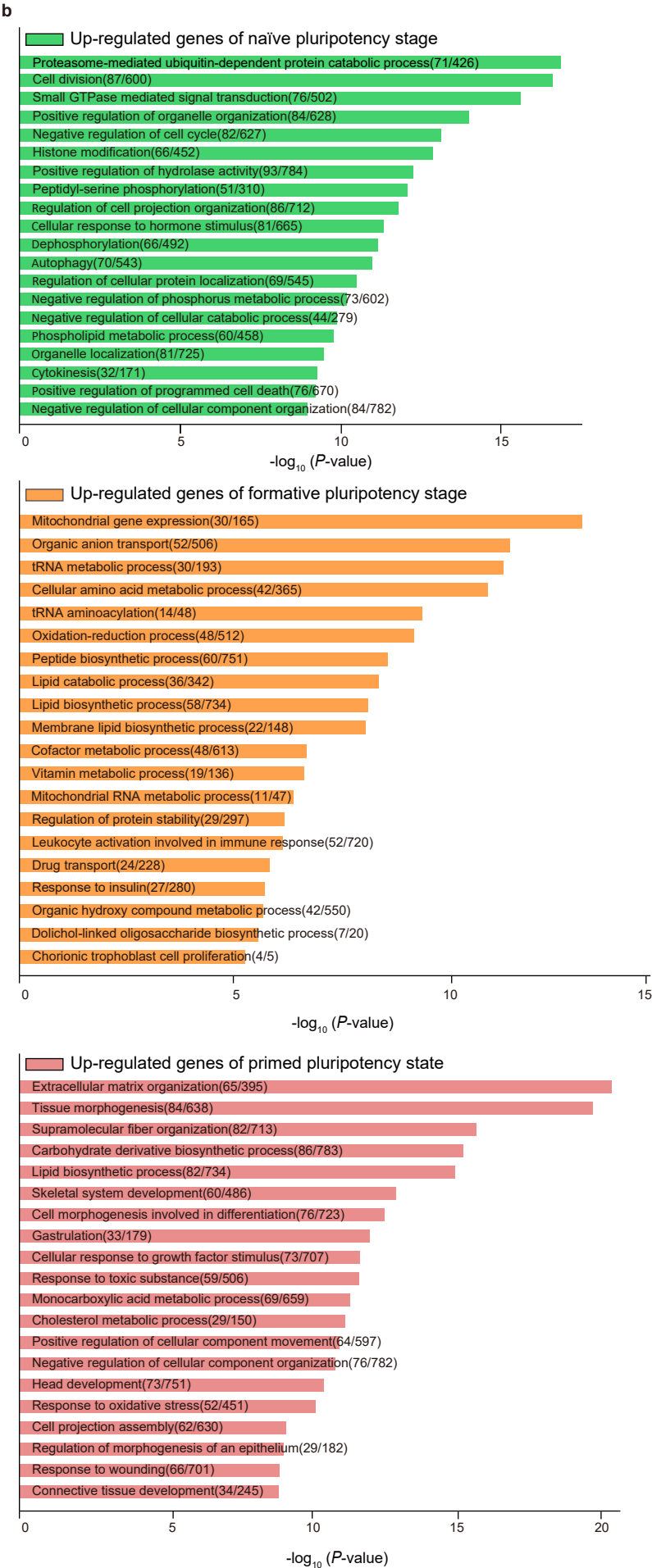
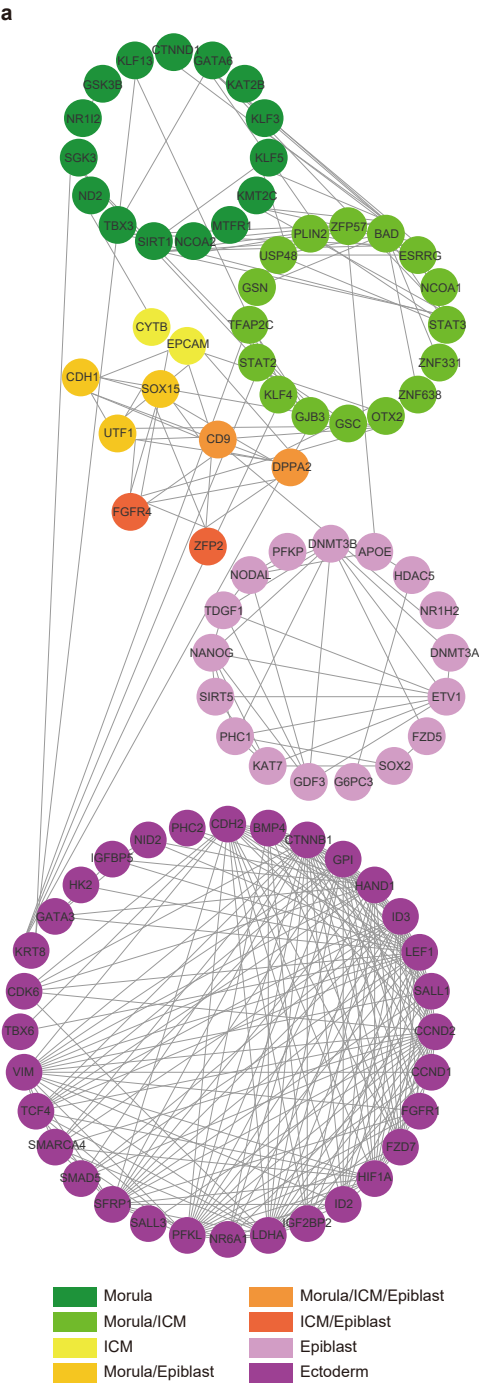


Fig. S2: Dynamic Changes of Pluripotency in Pig Pre-implantation Embryos, Related to Fig.

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a Network showing the differentially expressed genes (DEGs) in each cell type compared to other cell types. The different colors indicate different cell types. The circles denote the collections of DEGs. Each cell type is connected with its DEGs by the internal lines of the network. **b** A collection of four naïve pluripotency related clusters with functional enrichment for 1 735 genes from Fig. 1f (green bars), and 1 117 genes from four formative pluripotency related clusters (orange bars), and 1 289 genes from four primed pluripotency related clusters (red bars). The top functional terms of Metascape (<https://metascape.org>; see [Materials and methods](#)) summary gene set in each enriched cluster are shown, with the constraint of showing no more than 20 terms. The number after each term represents the hit genes out of total genes for this term.