## **SUPPLEMENTARY DATA**

## **Supplementary Table SI** Pregnancy outcomes and human oocyte trait quality assignment for weighted correlation network analysis.

Sample#	Age	YNG/AMA <sup>a</sup>	Maturation% <sup>b</sup>	Fertilization% <sup>c</sup>	GQ Blastocyst% <sup>d</sup>	Blastocyst% <sup>e</sup>	Birthsf
AMA GV/MII - I	40	l	I	0	Ι	Ι	2
AMA GV/MII - 2	43	1	0	0	0	0	0
AMA GV/MII - 3	40	1	1	0	0	0	0
AMA GV/MII - 4	42	1	1	1	1	1	0
AMA GV/MII - 5	43	1	1	1	NA	NA	0
YNG GV/MII - I	27	0	0	1	0	0	3
YNG GV/MII - 2	29	0	0	1	0	1	2
YNG GV/MII - 3	29	0	0	0	1	0	2
YNG GV/MII - 4	29	0	1	1	1	1	1
YNG GV/MII - 5	20	0	0	0	0	0	0

GV, germinal vesicle; MII, metaphase II.

A-E represent categorical values based on raw data. a: maternal age categorization - young (YNG, < 30 years) = 0, advanced maternal age (AMA,  $\geq$ 40 years) = 1; b: poor (<60%) = 0, moderate (65–72%) = 1, good (>80%) = 2; c: poor ( $\leq$ 75%) = 0, good (>75%) = 1; d: overall good quality blastocyst rate, poor ( $\leq$ 50%) = 0, good (>50%) = 1; e: overall blastocyst development rate of any grade, poor ( $\leq$ 70%) = 0, good (>70%) = 1. F represents the number of births from oocytes obtained during this cycle.