

Figure S1. Association of miR-205 expression with FIGO stage. (A) miR-205 across FIGO stages I-IV (Kruskal-Wallis test; $P=0.263$). (B) miR-205 in FIGO stages I-III combined vs. stage IV (Mann-Whitney U test; $P=0.067$). Relative expression, log₁₀ scale; boxes=IQR; central line=median; whiskers=1.5xIQR; points=individual samples. miR-205 tended to be higher in advanced (stage IV) disease, but the difference was not statistically significant. miR, microRNA; FIGO, International Federation of Gynecology and Obstetrics.

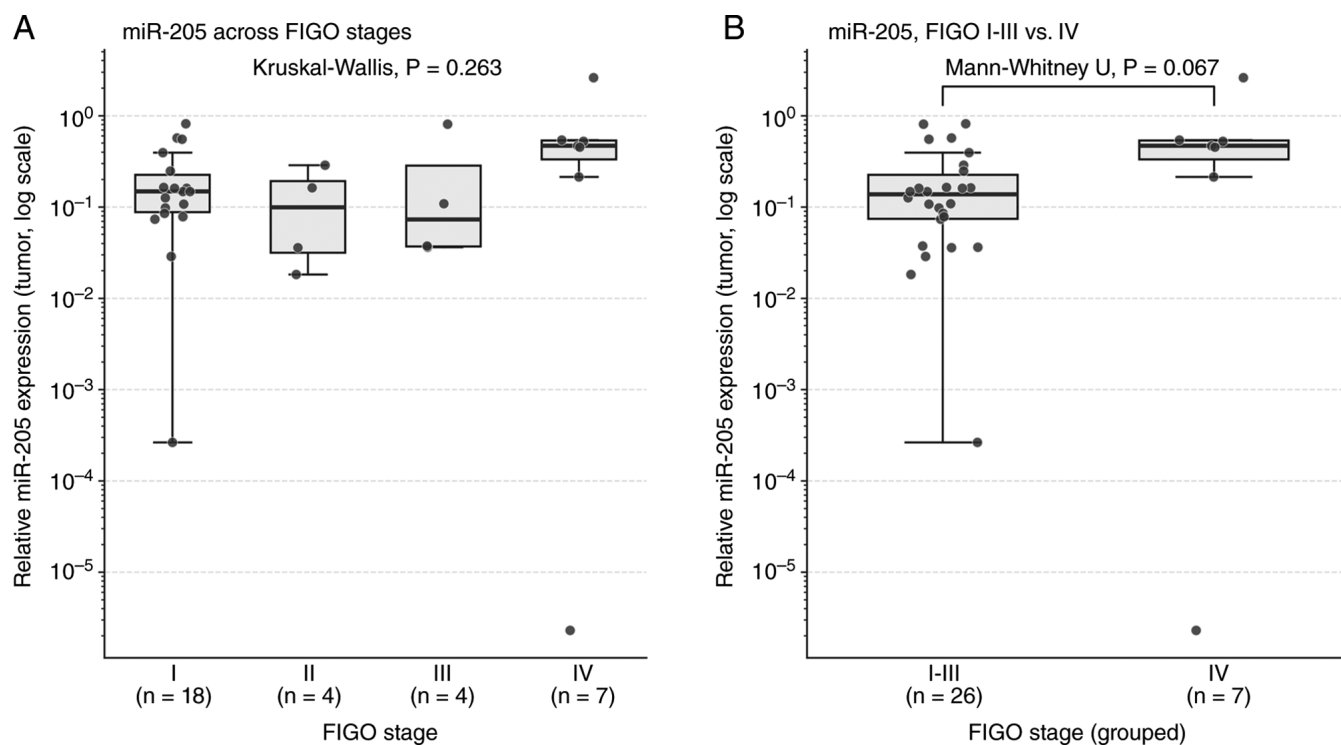


Figure S2. Association of miR-223 expression with FIGO stage (I-IV). Kruskal-Wallis test, $P=0.054$. Box-plot elements as in Fig. S1. No statistically significant association was observed miR, microRNA; FIGO, International Federation of Gynecology and Obstetrics.

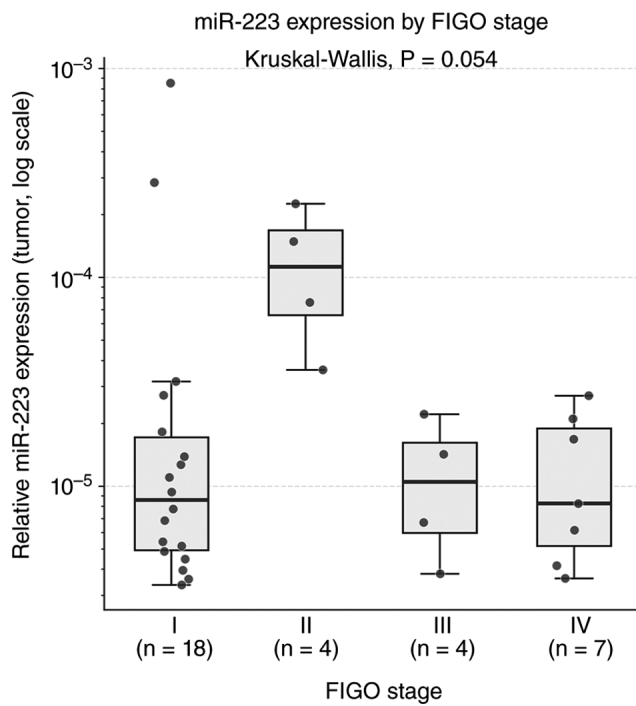


Figure S3. Association of miR-590 expression with FIGO stage (I-IV). Kruskal-Wallis test, $P=0.321$. Box-plot elements as in Fig. S1. No statistically significant association was observed. miR, microRNA; FIGO, International Federation of Gynecology and Obstetrics.

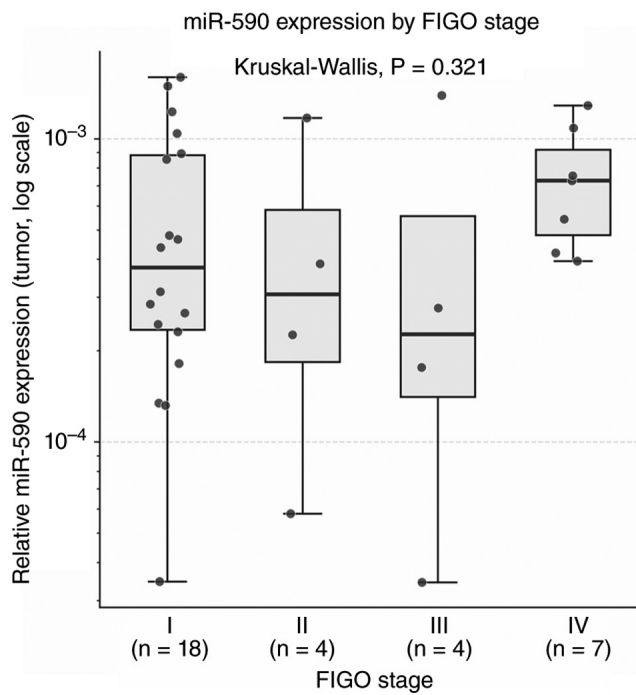


Figure S4. Association of miR-205 expression with tumor grade (G1, n=5; G2-3, n=29). Mann-Whitney U test, $P=0.393$. Box-plot elements as in Fig. S1. No statistically significant association was observed. miR, microRNA; FIGO, International Federation of Gynecology and Obstetrics.

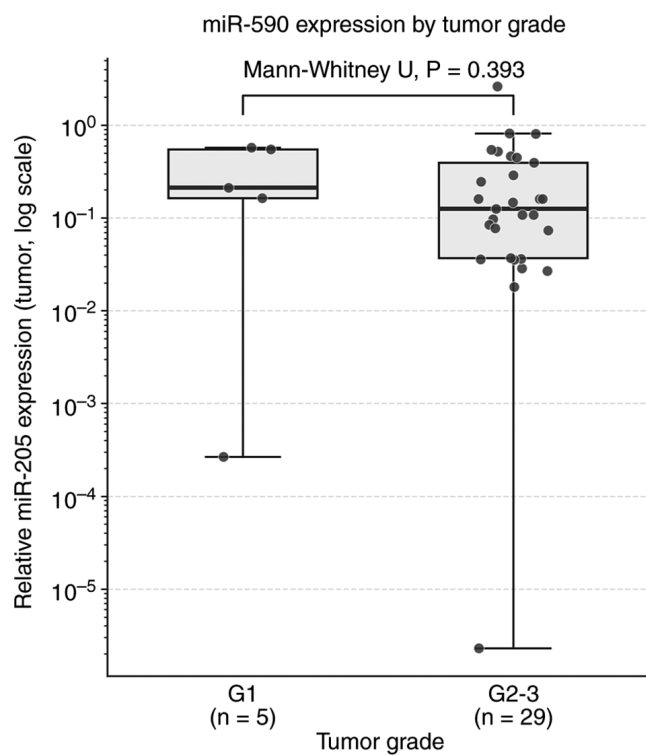


Figure S5. Association of miR-223 expression with tumor grade (G1, n=5; G2-3, n=29). Mann-Whitney U test, $P=0.851$. Box-plot elements as in Fig. S1. Although miR-223 expression appeared marginally higher in higher-grade tumors, this trend was not statistically significant. miR, microRNA; FIGO, International Federation of Gynecology and Obstetrics.

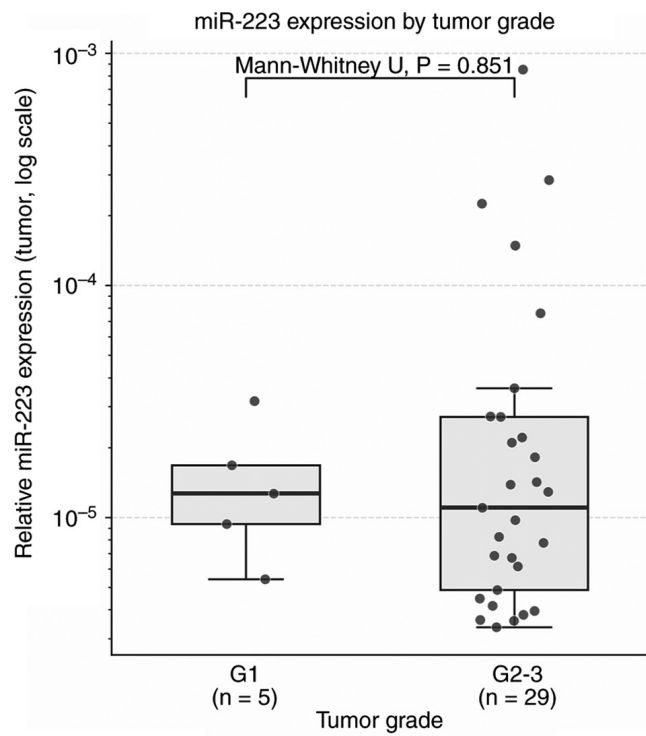


Figure S6. Overall survival according to miRNA fold-change (tumor/matched control) in the matched subset (n=20; 10 events), dichotomized at the median fold-change. (A) miR-205, (B) miR-223 and (C) miR-590. Vertical ticks denote censored observations; numbers at risk are shown below each panel. Each panel reports the log-rank P-value, the Cox hazard ratio (high vs. low fold-change) with 95% CI, and the 5-year (60-month) overall survival per group; the log-rank test was used for the primary comparison. Low miR-223 fold-change was associated with significantly improved OS (HR 6.18; 95% CI 1.24-30.79; log-rank P=0.013; 5-year OS 79% vs. 23%). miR-205 (log-rank P=0.592) and miR-590 (log-rank P=0.308) fold-change showed no significant association with OS. miR, microRNA; OS, overall survival; HR, hazard ratio; CI, confidence interval.

