

Supplementary Information:

Tracking the origin of synchronous endometrial and ovarian cancer by next-generation sequencing – a case report.

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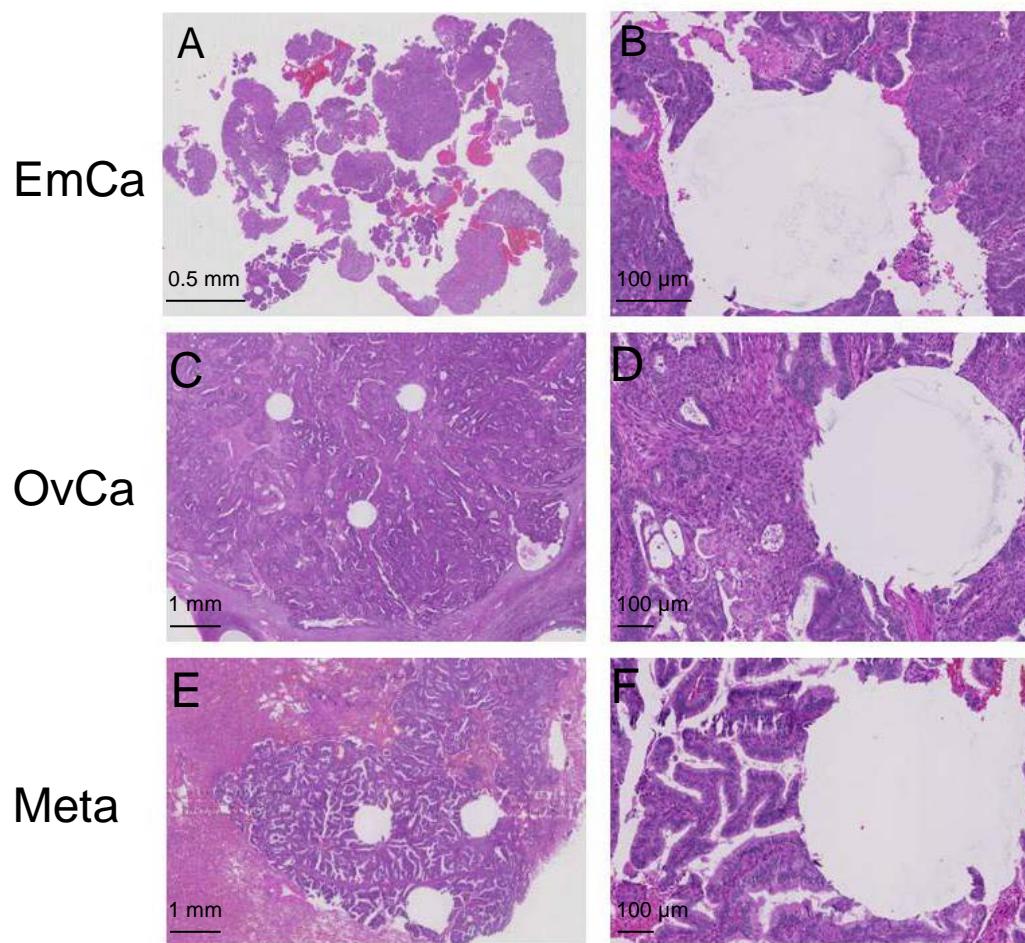
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Supplementary Figure 1A-F. Areas used for DNA extraction in the endometrial (A, B) and ovarian (C, D) carcinoma and the lung metastasis (E, F).

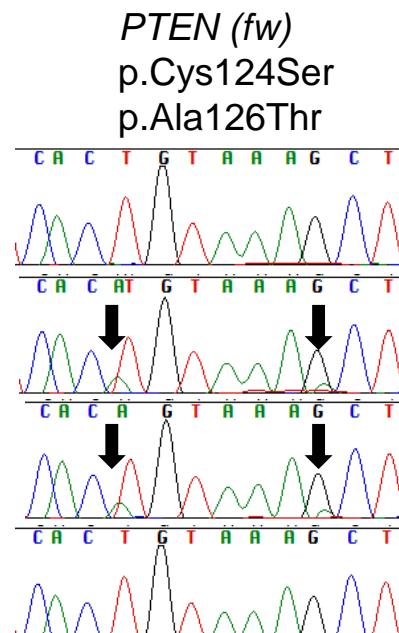
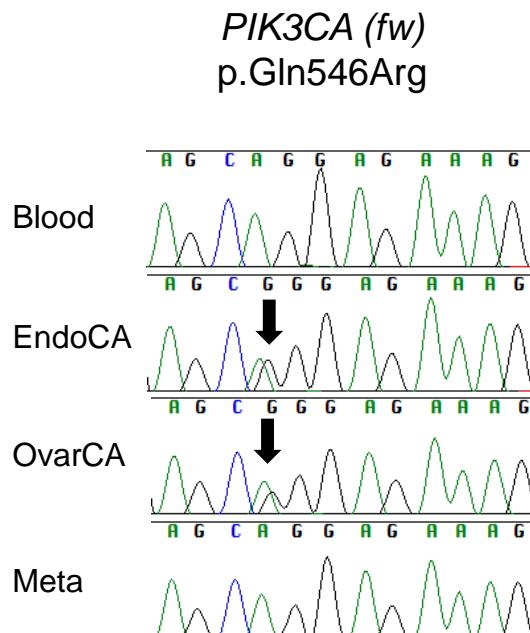
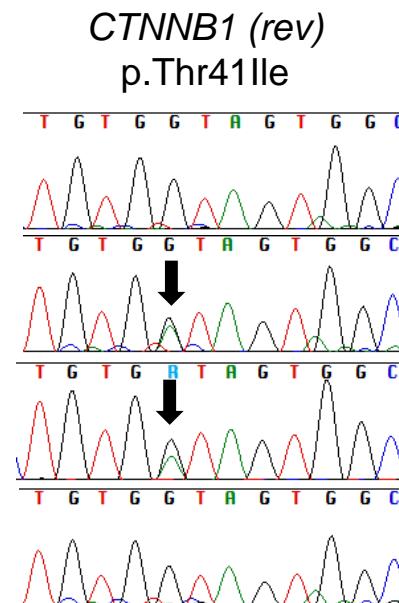
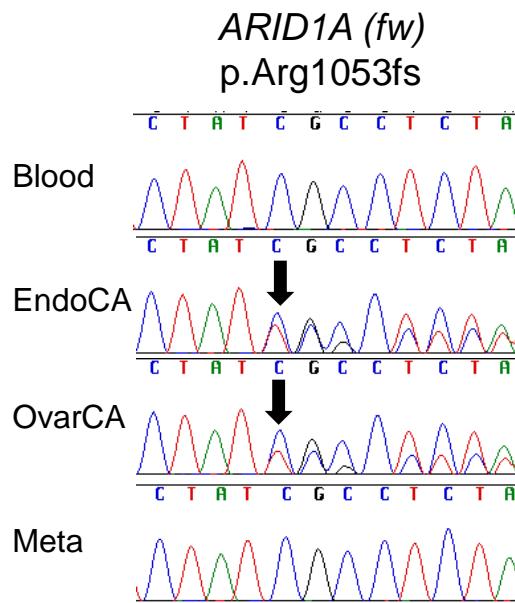
Supplementary Figure 2. Electropherograms from Sanger sequencing of *ARID1A*, *CTNNB1*, *PIK3CA* and *PTEN*. Arrows show the peaks of the mutated sites.

Suppl. Table 1. Sequences of the primers used for Sanger sequencing.

Suppl. Table 2. Comprehensive list of variants from targeted re-sequencing of the ovarian cancer, endometrial cancer and lung metastasis.



Suppl. Figure 1A-F. Areas used for DNA extraction in the endometrial (A, B) and ovarian (C, D) carcinoma and the lung metastasis (E, F). The oval gaps in the FFPE tissue correspond to the punches excised from the block and processed for NGS HE staining; EmCa, endometrial carcinoma; OvCa, ovarian carcinoma; Meta, lung metastasis.



Suppl. Figure 2. Electropherograms from Sanger sequencing of *ARID1A*, *CTNNB1*, *PIK3CA* and *PTEN*. Arrows show the peaks of the mutated sites.

Suppl. Table 1. Sequences of the primers used for Sanger sequencing.

Location	Gene	Forward primer	Reverse primer
chr1:27094447	ARID1A	TTTGACCTGAACCTTCCAGA	TCACCTGAGTCAATCCACCA
chr3:12641248	RAF1	ATGGGTTGATCCTTGATGC	ACACCAAAGCCCTGCAGTTA
chr3:41266125	CTNNB1	AAGCGGCTGTTAGTCAGTGG	TCAAAACTGCATTCTGACTTTCA
chr3:178936095	PIK3CA	CTGTGAATCCAGAGGGAAA	TTTAGCACTTACCTGTGACTCCA
chr6:152443737	SYNE1	GAAGGTGCAGAGGCAAAAG	GGGCACAGCTGTAGTCTTCC
chr7:106524631	PIK3CG	TGCTCTCTGAAAATGGTTCC	TCACCTGTCTCGGTGATCATA
chr10:89692886/92	PTEN	AGACCATAACCCACCACAGC	TGGTCCTTACTTCCCCATAGAA
chr20:40710566	PTPRT	AGGCCGTAGTGGAACCTTCT	TAGCAGCCAGGTGTTGAGTG
chrX:41057832	USP9X	TCCCTTTCAACTTTAGGAA	CCTCACACAGCCAACAGCTA

Suppl. Table 2. Comprehensive list of variants from targeted re-sequencing of blood, ovarian cancer, endometrial cancer and lung metastasis.