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| Figure S1. Unsupervised hierarchical clustering of differentially expressed miRNAs in HPV-related versus non-HPV-related PSCC (p ≤ 0.05; q ≥ 0.81; fold change ≥ 2). |

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| Figure S2. Heat map after unsupervised hierarchical clustering of differentially expressed miRNAs in tumor versus normal tissues (p ≤ 0.05; q ≥ 0.99; fold change ≥ 2). |

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| Figure S3. Heat map after unsupervised hierarchical clustering of differentially expressed miRNAs in metastatic versus non- metastatic (HPV-positive) basaloid, warty, and warty basaloid histological subtype (p ≤ 0.05; q ≥0.79; fold change ≥ 2). |

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| Figure S4. miRNAs without significant differences as determined by TaqMan® RT-qPCR (p ≤ 0.05). (a) and (b)Mean ΔCt value of (HPV-negative) usual PSCC and normal tissue samples; (c) mean ΔCt value of metastatic versus non-metastatic usual (HPV-negative) tumors; Lower ΔCt implays higher expression level. Mann-Whitney U test was used to determine p-values |