Supplementary Table S1. Histopathologic features and nucleic acid quality indices of excluded samples

| Variable | Sample 1 | Sample 2 | Sample 3 |
|--------------------------------|----------------------|---------------|---------------|
| Exclusion reason | Fibrotic tissue only | Mainly fibrin | Mainly fibrin |
| Organ | Liver | Kidney | Kidney |
| Year | 1 | 5 | 1 |
| Ischemic time (min) | 148 | 56 | 119 |
| Tumor cell proportion (%) | 0 | 0 | 0 |
| Histopathologic feature | | | |
| Inflammatory cell infiltration | Scanty | Scanty | Scanty |
| Stromal fibrosis | Marked | No | No |
| Necrosis (%) | 0 | 0 | 0 |
| Extracellular mucin pool (%) | 0 | 0 | 0 |
| DNA | | | |
| DNA integrity number | 7.2 | 7.6 | 2.7 |
| Concentration (ng/µL) | 59.429 | 32.442 | 35.798 |
| % Integrated area | 99.25 | 98.46 | 100.00 |
| RNA | | | |
| RNA integrity number | 2.8 | 0.0 | 2.4 |
| Concentration (ng/µL) | 8.184 | 2.171 | 3.237 |
| 28s/18s | 0.0 | 0.0 | 0.0 |

Histopathologic features and tapestation-based nucleic acid quality metrics of three samples excluded from downstream analysis due to absence of tumor cells.