

LoC-SERS platform based on triple signal amplification for highly sensitive detection of colorectal cancer miRNAs

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Table S1. Oligonucleotides used in the experiments

| Name | Sequence (5' - 3') |
|-----------|--|
| miRNA-122 | UGG AGU GUG ACA AUG GUG UUU G |
| miRNA-192 | CUG ACC UAU GAA UUG ACA GCC |
| MT1-1 | UGG ACU GUG ACA AUG GUG UUU G |
| MT1-2 | CGG ACC UAU GAA UUG ACA GCC |
| MT3-1 | UCG AGU GUG AUA ACG GUG UUU G |
| MT3-2 | CAG ACC UGU GAA UGG ACA GCC |
| hpDNA1-1 | ATG TGT TGC AGT CGA TGC TGA GGC AAA CAC CAT TGT CAC GTA GTG CAA CA |
| hpDNA1-2 | TCA GAT GTG ACT AGT GCT AGC TGA GGC TGT CAA TTC ATA CAT GTC ACA T |
| hpDNA2-1 | ACG TAA CAC CAT TGT CAC GAC CAT CGA CTT ATC GTG ACA ATG GTG TTG CCT CA |
| hpDNA2-2 | ACG GTC AAT TCA TAC ATG TAA TAG ACA TGC TAC ATG TAT GAA TTG ACA GCC TCA |
| hpDNA3-1 | CTA ACC ATC GAC TTA TAA CAC CAT TGT CAC GAT AAG TCG ATG GTC GTG ACA ACG C |
| hpDNA3-1 | CTA ACC ATC GAC TTA TAA CAC CAT TGT CAC GAT AAG TCG ATG GTC GTG ACA ACG C |
| DNA1-1 | ATG TGT TGC AGT CGA TGC |
| DNA1-2 | TCA GAT GTG ACT AGT GCT AGC |
| DNA2-1 | TGA GGC AAA CAC CAT TGT CAC GTA GTG CAA CA |
| DNA2-2 | TGA GGC TGT CAA TTC ATA CAT GTC ACA T |

Table S2 Basic characteristics of the healthy subjects and colorectal cancer patients

| Groups | Healthy subjects | Colorectal cancer |
|---------------|-------------------------|--------------------------|
| Average age | 34 | 47 |
| Gender | | |
| Male | 16 | 18 |
| Female | 14 | 12 |
| Sample number | 30 | 30 |

Table S3 The miR-122 and miR-192 concentration detected by SERS using the proposed platform and ELISA in colorectal cancer patients

| Samples | SERS (fM) | | qRT-PCR (fM) | |
|---------|-----------|---------|--------------|---------|
| | miR-122 | miR-192 | miR-122 | miR-192 |
| 1 | 28.65 | 18.60 | 27.18 | 19.61 |
| 2 | 24.78 | 19.67 | 29.68 | 19.01 |
| 3 | 26.99 | 23.77 | 22.40 | 20.43 |
| 4 | 22.51 | 29.40 | 24.20 | 28.29 |
| 5 | 22.09 | 24.46 | 26.08 | 24.55 |
| 6 | 29.71 | 26.06 | 28.99 | 25.15 |
| 7 | 27.37 | 29.65 | 30.63 | 25.25 |
| 8 | 22.31 | 25.10 | 29.33 | 22.87 |
| 9 | 31.96 | 27.60 | 26.49 | 21.39 |
| 10 | 21.88 | 28.97 | 21.80 | 20.48 |
| 11 | 26.23 | 24.62 | 23.81 | 25.58 |
| 12 | 30.86 | 24.05 | 27.01 | 24.28 |
| 13 | 29.93 | 25.95 | 25.58 | 21.23 |
| 14 | 28.52 | 24.54 | 25.95 | 28.90 |
| 15 | 30.94 | 22.57 | 29.38 | 28.98 |
| 16 | 32.99 | 24.36 | 31.97 | 21.43 |
| 17 | 33.41 | 25.60 | 30.55 | 24.30 |
| 18 | 32.24 | 29.07 | 28.53 | 27.47 |
| 19 | 26.03 | 26.29 | 28.36 | 23.91 |
| 20 | 28.54 | 20.61 | 22.96 | 21.42 |
| 21 | 27.84 | 20.86 | 30.07 | 18.63 |
| 22 | 30.36 | 21.02 | 27.43 | 21.66 |
| 23 | 24.67 | 31.40 | 21.69 | 29.12 |
| 24 | 24.21 | 27.74 | 23.65 | 25.11 |
| 25 | 27.89 | 21.08 | 25.20 | 18.98 |
| 26 | 26.15 | 27.66 | 26.33 | 23.53 |
| 27 | 26.91 | 28.00 | 28.97 | 28.81 |
| 28 | 32.20 | 25.50 | 32.32 | 22.20 |
| 29 | 26.07 | 31.33 | 24.68 | 28.18 |
| 30 | 30.77 | 25.75 | 22.42 | 23.48 |

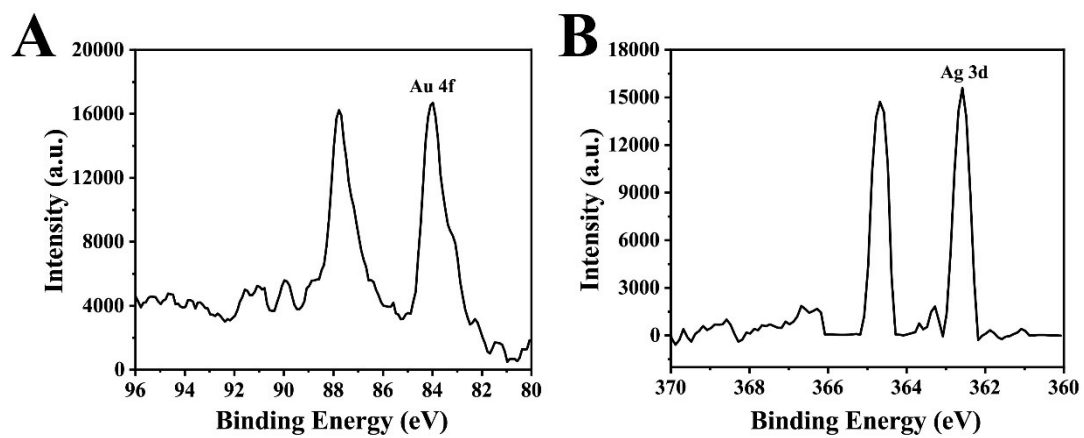


Fig. S1. XPS spectra of Au-AgNSs (a) high resolution Au 4f (b) high resolution Ag 3d.