**Additional file 1**

**Additional file 1: Table S1 Primer sequences and target sequence.**

|  |  |
| --- | --- |
| **Name** | **Forward** |
| **18S forward** | CGGCTACCACATCCAAGGAA |
| **18S reverse** | GCTGGAATTACCGCGGCT |
| **SNAP23 forward** | CTGCCCATGTAATAGAACAAAGAAC |
| **SNAP23 reverse** | CGGGCCTGGCTGTTTAGATA |
| **TFAM forward** | CGCTCCCCCTTCAGTTTTGT |
| **TFAM reverse** | CCAACGCTGGGCAATTCTTC |
| **CREB1 forward** | ACCAAGTTGTTGTTCAAGGTACT |
| **CREB1 reverse** | ACATGTTACCATCTTCAAACTGACG |
| **Lin28a forward** | CCAGTGGATGTCTTTGTGCACC |
| **Lin28a reverse** | GTGACACGGATGGATTCCAGAC |
| **Lin28b forward** | CCTGTTTAGGAAGTGAAAGAAGAC |
| **Lin28b reverse** | CCTGTTTAGGAAGTGAAAGAAGAC |
| **VAMP3 forward** | GCTCTCTGAGTTAGACGACCGT |
| **VAMP3 reverse** | CCAGAACAGTAATCCCGATTGCC |
| **Rab35 forward** | CAGCCCATCTTACTGCAAGCAG |
| **Rab35 reverse** | GCTGACAACCTGTCGGAGAGAA |
| **U6 forward** | GCTCGCTTCGGCAGCACATATAC |
| **U6 reverse** | GTCGTATCGCTCGCTTCGGCAGCACATATACCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACAAAATATGG |
| **Universal-reverse** | AGTGCAGGGTCCGAGGTATT |
| **Let-7a forward** | GCGCGTGAGGTAGTAGGTTGT |
| **Let-7a reverse** | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACAACTAT |
| **sh-SNAP23-1 forward** | GCTTGGACCAAATAAATAA |
| **sh-SNAP23-1 reverse** | GCTTGGACCAAATAAATAA |
| **sh-SNAP23-2 reverse** | GCAGTATCCTGGGAAATCT |
| **sh-SNAP23-2 forward** | GCAGTATCCTGGGAAATCT |
| **sh-SNAP23-3 reverse** | GCCACAACGAAGATCTAGT |
| **sh-SNAP23-3 forward** | GCCACAACGAAGATCTAGT |
| **SNAP23 vector reverse** | ggatccTTACTTGTCGTCATCGTCTTTGTAGTCGCTGTCAATGAGTTTCTTTGCT |
| **SNAP23 vector forward** | gaattcGCCACCATGGATAATCTGTCATCAGAAGAA |

**Additional file 1: Table S2 Comparison of clinical pathologic characteristics of let-7a in serum EVs from CRC patient.**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Variable | n | Low | High | Pearson x | p Value |
| Total | 31 | 15 | 16 |  |  |
|  | Gender |
| MaleFemale | 1813 | 10 55.6%5 38.5% | 8 44.4%8 61.5% | 0.883 | 0.347 |
|  | Age |
| <65>=65 | 724 | 5 71.4%10 41.7% | 2 28.6%14 58.3% | 1.922 | 0.166 |
|  | Distant metastasis  |
| NoYes | 256 | 12 48.0%3 50.0% | 13 52.0%3 50.0% | 0.008 | 0.930 |
|  | AJCC stage |
| 1-23-4 | 922 | 6 66.7%9 40.9% | 3 33.3%13 59.1% | 1.697 | 0.193 |

**Additional file 1: Table S3 Comparison of clinical pathologic characteristics of CRC patient tissue with SNAP23 expression.**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Variable | n | Weak | Moderate | Strong | Pearson x | p Value |
| Total | 31 | 7 | 13 | 11 |  |  |
|  | Gender |
| MaleFemale | 1813 | 4 22.2%3 23.1% | 6 33.3%7 53.8% | 8 44.4%3 23.1% | 1.731 | 0.421 |
|  | Age |
| <65>=65 | 724 | 3 42.9%4 16.7% | 2 28.6%11 45.8% | 2 28.6%6 37.5% | 2.153 | 0.341 |
|  | Distant metastasis  |
| NoYes | 256 | 6 24.0%1 16.7% | 10 40.0%3 50.0% | 9 36.0%2 33.3% | 0.240 | 0.887 |
|  | AJCC stage |
| 1-23-4 | 922 | 4 44.4%3 13.6% | 2 22.2%11 50.0% | 3 33.3%8 36.4% | 3.876 | 0.144 |