Supplementary Figure 2

(A) Percent survival of different proteins:

- **ACE2**
  - Survival time (Days)
  - Number at risk: high 114, low 102
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 0.545 (0.853-1.087)
  - pValue: 0.18

- **DPP4**
  - Survival time (Days)
  - Number at risk: high 164, low 52
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 0.883 (0.783-0.996)
  - pValue: 0.17

- **ANPEP**
  - Survival time (Days)
  - Number at risk: high 180, low 36
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 1.243 (1.094-1.413)
  - pValue: 0.043

- **AXL**
  - Survival time (Days)
  - Number at risk: high 140, low 76
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 1.227 (1.006-1.497)
  - pValue: 0.11

- **TMPRSS2**
  - Survival time (Days)
  - Number at risk: high 56, low 160
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 0.963 (0.853-1.087)
  - pValue: 0.17

- **ENPEP**
  - Survival time (Days)
  - Number at risk: high 124, low 92
  - Percent survival:
    - high: 0.75, 0.50, 0.25, 0.00
    - low: 0.75, 0.50, 0.25, 0.00
  - Hazard ratio: 0.567 (0.444-0.725)
  - pValue: 0.00086

(B) P-values and Hazard ratios:

| Variable          | pValue  | Hazard ratio          |
|-------------------|---------|-----------------------|
| ANPEP             | 0.545   | 0.963 (0.853-1.087)   |
| DPP4              | 0.043   | 0.883 (0.783-0.996)   |
| ENPEP             | <0.001  | 1.243 (1.094-1.413)   |
| PRS_type          | <0.001  | 1.974 (1.682-2.317)   |
| Histology         | 0.103   | 0.692 (0.445-1.077)   |
| Grade             | <0.001  | 2.688 (1.965-3.677)   |
| Age               | 0.043   | 1.227 (1.006-1.497)   |
| Chemo             | <0.001  | 0.660 (0.520-0.838)   |
| IDH_mutation      | <0.001  | 0.567 (0.444-0.725)   |
| 1p19q_codeletion  | <0.001  | 0.387 (0.278-0.540)   |