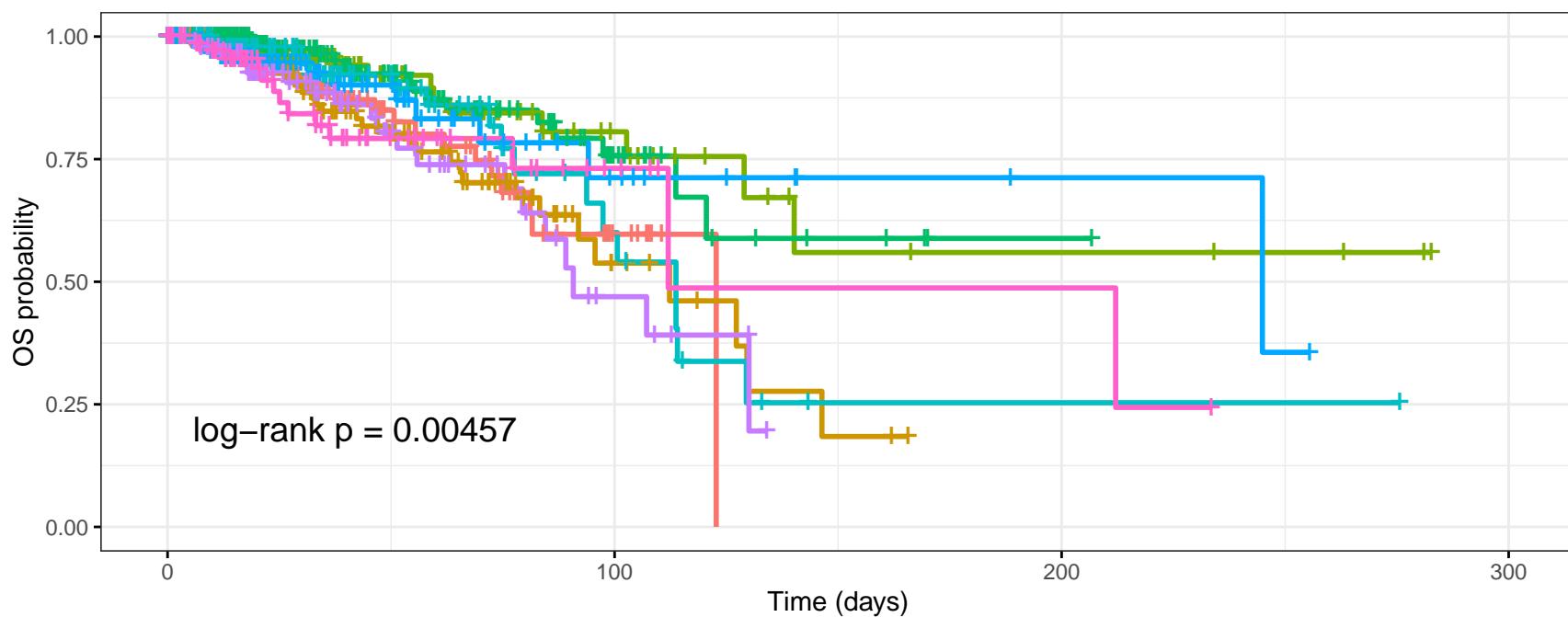


TNF signaling pathway – Expr quartiles x CNA (8 groups, OS, TCGA–BRCA)

Quartiles of expression (Q1–Q4) crossed with CNA (0/1)

- + Q1 (lowest) & CNA = 0 + Q3 & CNA = 0 + Q1 (lowest) & CNA = 1 + Q3 & CNA = 1
- + Q2 & CNA = 0 + Q4 (highest) & CNA = 0 + Q2 & CNA = 1 + Q4 (highest) & CNA = 1



Number at risk

| | | | | |
|------------------------|-----|----|---|---|
| Q1 (lowest) & CNA = 0 | 148 | 7 | 0 | 0 |
| Q2 & CNA = 0 | 150 | 9 | 0 | 0 |
| Q3 & CNA = 0 | 159 | 16 | 4 | 0 |
| Q4 (highest) & CNA = 0 | 184 | 16 | 1 | 0 |
| Q1 (lowest) & CNA = 1 | 111 | 10 | 1 | 0 |
| Q2 & CNA = 1 | 108 | 9 | 2 | 0 |
| Q3 & CNA = 1 | 99 | 6 | 0 | 0 |
| Q4 (highest) & CNA = 1 | 74 | 6 | 2 | 0 |