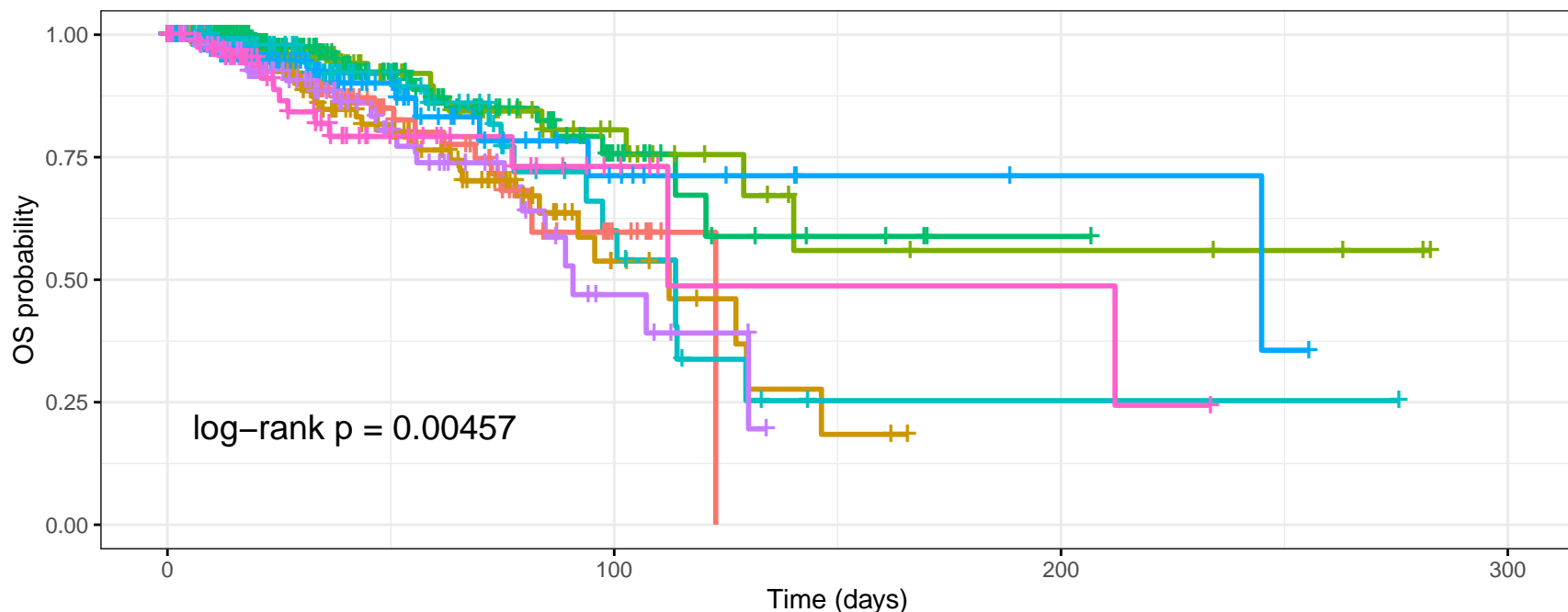
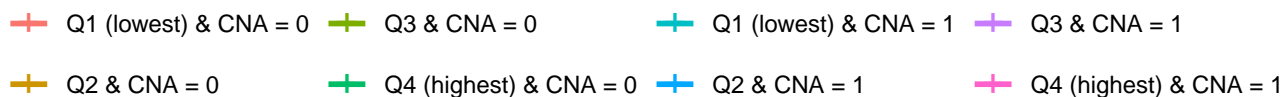


# TNF signaling pathway – Expr quartiles x CNA (8 groups, OS, TCGA–BRCA)

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



## 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   |
|                        | 0   | 100 | 200 | 300 |