

**Supplemental Digital Content 5. Initial cohort: nominal associations of SNPs with overall survival (OS, p<0.05) in Cox regression models adjusted for stage.** Results are ranked by significance of OS-SNP associations. The associations with p>0.05 are available upon request.

| SNP        | Alleles | Gene         | HR   | 95% CI       | P value | Q value |
|------------|---------|--------------|------|--------------|---------|---------|
| rs7996030  | G>A     | <i>FLT1</i>  | 1.59 | 1.15 to 2.21 | 0.0052  | 0.0848  |
| rs9582036  | A>C     | <i>FLT1</i>  | 1.52 | 1.13 to 2.04 | 0.0054  | 0.0848  |
| rs12813551 | T>C     | <i>KRAS</i>  | 0.65 | 0.47 to 0.90 | 0.0089  | 0.0937  |
| rs10842513 | C>T     | <i>KRAS</i>  | 1.53 | 1.08 to 2.17 | 0.0168  | 0.1111  |
| rs10505980 | G>A     | <i>KRAS</i>  | 0.68 | 0.49 to 0.93 | 0.0176  | 0.1111  |
| rs542403   | A>G     | <i>FRS2</i>  | 0.63 | 0.41 to 0.95 | 0.0294  | 0.1544  |
| rs11549465 | C>T     | <i>HIF1A</i> | 1.61 | 1.01 to 2.56 | 0.0437  | 0.1554  |
| rs7219     | A>G     | <i>GRB2</i>  | 0.65 | 0.43 to 0.99 | 0.0471  | 0.1554  |
| rs1570360  | G>A     | <i>VEGFA</i> | 1.37 | 1.00 to 1.88 | 0.0496  | 0.1554  |