

Supplementary Table 2: Univariate analysis in the 238 tumors set from Fan *et al.*

| | TRIM24 mRNA levels | | |
|-----------------------|--------------------|--------------|--------|
| | HR | 95%CI | P* |
| TRIM24 | | | |
| ≤ 1.08 | 1 | | <0.001 |
| > 1.08 | 2.68 | [1.5 ; 4.8] | |
| N | | | |
| N0 | 1 | | 0.414 |
| N+ | 0.82 | [0.5 ; 1.3] | |
| T | | | |
| T1 (≤2 cm) | 1 | | 0.003 |
| T2-T3-T4 (>2cm) | 2.1 | [1.3 ; 3.4] | |
| ER | | | |
| ER- | 1 | | <0.001 |
| ER+ | 0.3 | [0.2 ; 0.5] | |
| Grade SBR | | | |
| I | 1 | | <0.001 |
| II | 4.9 | [1.5 ; 16] | |
| III | 12.1 | [3.7 ; 38.9] | |
| Type / Sorlie | | | |
| Normal Like | 1 | | <0.001 |
| Luminal A | 0.64 | [0.2 ; 1.8] | |
| Luminal B | 2.39 | [0.9 ; 6.5] | |
| ErbB2 | 2.65 | [0.9 ; 7.5] | |
| Basal | 3.52 | [1.3 ; 9.3] | |
| 70-gene | | | |
| Good | 1 | | <0.001 |
| Poor | 10.8 | [4.3 ; 27] | |
| Two-gene ratio | | | |
| High | 1 | | 0.028 |
| Low | 0.57 | [0.3 ; 0.9] | |
| Wound response | | | |
| Activated | 1 | | <0.001 |
| Quiescent | 0.19 | [0.07 ; 0.5] | |
| Recurrence | | | |
| High | 1 | | <0.001 |
| Intermediate | 0.27 | [0.1 ; 0.7] | |
| Low | 0.06 | [0.01 ; 0.2] | |

* Logrank test