|  | 2016 BRAF Wild-type | | 2016 BRAF Mutant | | 2018 BRAF Wild-type | |
| --- | --- | --- | --- | --- | --- | --- |
|  | HR (95% CI)1 | p-value | HR (95% CI)1 | p-value | HR (95% CI)1 | p-value |
| **BRAF status - Year of diagnosis** |  |  |  |  |  |  |
| *2016 BRAF WT* | — |  | 0.93 (0.72 to 1.20) | 0.57 | 1.23 (0.94 to 1.61) | 0.13 |
| *2016 BRAF MT* | 1.08 (0.83 to 1.40) | 0.57 | — |  | 1.33 (1.01 to 1.74) | **0.041** |
| *2018 BRAF WT* | 0.81 (0.62 to 1.06) | 0.13 | 0.75 (0.57 to 0.99) | **0.041** | — |  |
| *2018 BRAF MT* | 0.79 (0.61 to 1.03) | 0.085 | 0.74 (0.56 to 0.96) | **0.024** | 0.98 (0.74 to 1.29) | 0.87 |
| 1HR = Hazard Ratio, CI = Confidence Interval | | | | | | |