**Supplemental Table 1.** Univariate binomial logistic regression models for VT morphology prediction (all RBBB-VT, n=52, compared to LBBB-VT, n=486)

|  |  |
| --- | --- |
| VT morphology | RBBB VT (n=52) |
|  | Odds Ratio | CI 95% | CI 95% | p value |
| Female | 1.52 | 0.79 | 2.8 | 0.190 |
| Age at first VT | 1.01 | 0.99 | 1.03 | 0.434 |
| *PKP2* mutation | 0.25 | 0.11 | 0.51 | **<0.001** |
| *DSG2* mutation | 1.13 | 0.26 | 3.37 | 0.847 |
| *DSP* mutation | 6.98 | 3.01 | 15.61 | **<0.001** |
| *JUP* mutation | 0.93 | 0.05 | 5.02 | 0.948 |
| *DSC2* mutation | 1.04 | 0.06 | 5.7 | 0.971 |
| *PLN* mutation | 3.87 | 1.2 | 10.77 | **0.014** |