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# Prevalence and Prognostic Impact of Pathogenic Variants in Patients With Dilated Cardiomyopathy Referred for Ventricular Tachycardia Ablation

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## Abstract

**Objectives:** This study aimed to assess the frequency of (likely) pathogenic variants (LP/Pv) among dilated cardiomyopathy (DCM) ventricular tachycardia (VT) patients referred for CA and their impact on procedural outcome and long-term prognosis.

**Background:** The prevalence of genetic variants associated with monomorphic VT among DCM is unknown.

**Methods:** Ninety-eight consecutive patients (age  $56 \pm 15$  years; 84% men, left ventricular ejection fraction [LVEF]  $39 \pm 12\%$ ) referred for DCM-VT ablation were included. Patients underwent electroanatomical mapping and testing of  $\geq 55$  cardiomyopathy-related genes. Mapping data were analyzed for low-voltage areas and abnormal potentials. LP/Pv-positive (LP/Pv+) patients were compared with LP/Pv-negative (LP/Pv-) patients and followed for VT recurrence and mortality.

**Results:** In 37 (38%) patients, LP/Pv were identified, most frequently LMNA (n = 11 of 37, [30%]), TTN (n = 6 of 37, [16%]), PLN (n = 6 of 37, [16%]), SCN5A (n = 3 of 37, [8%]), RBM20 (n = 2 of 37, [5%]) and DSP (n = 2 of 37, [5%]). LP/Pv+ carriers had lower LVEF ( $35 \pm 13\%$  vs. LP/Pv-:  $42 \pm 11\%$ ;  $p = 0.005$ ) and were less often men (n = 27 [73%] vs. n = 55 [90%];  $p = 0.03$ ). After a median follow-up of 2.4 years (interquartile range: 0.9 to 4.4 years), 63 (64%) patients had VT recurrence (LP/Pv+: 30 of 37 [81%] vs. LP/Pv-: 33 of 61 [54%];  $p = 0.007$ ). Twenty-eight patients (29%) died (LP/Pv+: 19 of 37 [51%] vs. LP/Pv-: 9 of 61 [15%];  $p < 0.001$ ). The cumulative 2-year VT-free survival was 41% in the total cohort (LP/Pv+: 16% vs. LP/Pv-: 54%;  $p = 0.001$ ). The presence of LP/Pv (hazard ratio: 1.9; 95% confidence interval: 1.1 to 3.4;  $p = 0.02$ ) and unipolar low-voltage area size/cm<sup>2</sup> increase (hazard ratio: 2.5; 95% confidence interval: 1.6 to 4.0;  $p < 0.001$ ) were associated with a decreased 2-year VT-free survival.

**Conclusions:** In patients with DCM-VT, a genetic cause is frequently identified. LP/Pv+ patients have a lower LVEF and more extensive VT substrates, which, in combination with disease progression, may contribute to the poor prognosis. Genetic testing in patients with DCM-VT should therefore be recommended.

**Keywords:** catheter ablation; dilated cardiomyopathy; genetic mutation; genetic testing; genetic variant; inherited cardiomyopathy; nonischemic cardiomyopathy; ventricular tachycardia.

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