

1           **Recovery of cardiac function in cardiomyopathy due to titin truncation**

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25 **Introduction:** Dilated cardiomyopathy (DCM) is a frequent cause of heart failure and  
26 a common indication for heart transplantation. DCM has a strong genetic basis and  
27 the most common disease-causing mutations are variants that truncate the  
28 sarcomeric protein titin (TTN truncating variants (TTNtrvs); prevalence in familial or  
29 idiopathic DCM 25%<sup>1</sup> and 13%<sup>2</sup>, respectively). The prognosis of DCM is poor, but  
30 functional recovery from end-stage failure has been reported following both optimal  
31 medical therapy<sup>3</sup> and left ventricular assist device (LVAD) support<sup>4,5</sup>, though the  
32 determinants of successful recovery are unknown. It has been proposed that  
33 recovery from genetic cardiomyopathy may not be expected since the underlying  
34 cause is irreversible; whereas recovery may be more likely when DCM is due to  
35 reversible, non-genetic factors (e.g. myocarditis)<sup>6</sup>. To address this directly, we  
36 sequenced TTN in end-stage DCM patients who either recovered or did not recover  
37 following LVAD support.

38

39 **Methods:** We sequenced TTN in 70 cases referred to the Royal Brompton and  
40 Harefield National Health Service Trust between 1998 and 2010 for LVAD  
41 implantation due to non-ischaemic, medically refractory, end-stage DCM. Of these,  
42 29 patients recovered cardiac function during LVAD support and had their LVAD  
43 explanted. The other 41 patients did not recover cardiac function and were  
44 transplanted or died while on LVAD support. A pharmacological regime designed to  
45 promote recovery (combination therapy<sup>4,5</sup>) was used in 35/70 patients and  
46 continued after explantation. The study was approved by institutional ethics  
47 committees, with written informed consent from participants. Targeted next-  
48 generation sequencing (NGS) was performed using an assay designed to assess all

49 known coding exons in TTN<sup>2</sup>. Genetic variants in NGS data were identified as  
50 previously described<sup>2</sup> and were confirmed independently. Statistical comparisons  
51 between groups were tested using Fisher's exact test, ANOVA and unpaired t-test as  
52 appropriate. Differences in survival rates were tested using the Mantel-Cox test.  
53 Statistical significance was defined as a P-value of <0.05.

54

55 **Results:** We identified TTNtvs in 10 out of 70 cases (14% of total; Table 1). All TTNtvs  
56 were either novel or very rare<sup>α</sup> and located in exons constitutively expressed in the  
57 heart and, as such, considered disease-causing<sup>2</sup>. Of the patients with a TTNtv, 6/10  
58 recovered sufficient cardiac function to enable LVAD explantation. There was no  
59 statistical difference in TTNtv frequency between recovery cases and those who  
60 were transplanted or died on the device (6/29 [21%] vs. 4/41 [10%] respectively,  
61 P=0.30), and no evidence of clinical differences between TTNtv-positive and TTNtv-  
62 negative cases at the time of LVAD implantation (Table 1). Comparing the transplant-  
63 free survival rate in recovered patients, we found no difference between TTNtv-  
64 positive and TTNtv-negative cases; at three years post-explant, 4/6 (67%) TTNtv-  
65 positive cases were free from death and transplantation compared to 17/23 (74%)  
66 TTNtv-negative cases (Figure 1; P=0.74).

67

68 **Discussion:** Sustained improvement in cardiac function is observed in end-stage  
69 DCM following medical therapy and LVAD support, but it was previously unknown  
70 whether recovery could be achieved in DCM due to a genetic cause. Here, we show  
71 that recovery is possible in DCM due to a truncating mutation in the TTN gene. We  
72 also present the preliminary findings that DCM with a TTNtv is as equally recoverable

73 as DCM without a TTNtv and that the long-term durability of recovery is also  
74 comparable. These observations now require replication in multi-centre prospective  
75 studies. Since TTNtvs are the most common genetic cause of DCM these results have  
76 important implications for patient selection for recovery programs.

77 "Nine TTNtvs were not present in ExAC, whilst one variant had a minor allele  
78 frequency of 0.0000166 (<http://exac.broadinstitute.org/>).

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80 **Author contributions**

81 Dr Felkin had full access to all of the data in the study and takes responsibility for the  
82 integrity of the data and the accuracy of the data analysis.

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89 The funders had no role in the design and conduct of the study; collection,  
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92 **Additional Contributions**

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124 **Table 1:** TTNtv status and clinical features of LVAD-supported, end-stage DCM  
125 patients who either recovered cardiac function and were successfully explanted  
126 (recovered), or who were transplanted or died with the device *in situ* (not  
127 recovered).

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129

130 **Figure 1. TTNtv and survival in recovered DCM patients.** In the three years  
131 following successful LVAD explanation, the actuarial rate of survival and freedom  
132 from transplantation at 1, 2 and 3 years post-explant in TTNtv-positive cases was  
133 83%, 83% and 67%. In TTNtv-negative cases, the rate was 88%, 88% and 75%.  
134 Differences between the survival rates were tested using the Mantel-Cox test.

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Variable	TTNtv, n=10		No TTNtv, n=60		P value
	Recovered	Not recovered	Recovered	Not recovered	
Number of patients	6	4	23	37	0.30 <sup>A</sup>
Male, n (%)	6 (100)	4 (100)	18 (78)	30 (81)	0.29 <sup>A</sup>
Clinical comments	-	1 post-chemo	2 PPCM	3 PPCM 2 post-chemo	
Family history, n (%)	2 (33)	0 (0)	2 (8.7)	1 (2.7)	1.0 <sup>A</sup>
Survived > 30 days post LVAD implant, n (%)	6 (100)	2 (50)	23 (100)	34 (92)	0.13 <sup>A</sup>
Received combination therapy <sup>3</sup> , n (%)	6 (100)	2 (50)	23 (100)	14 (38)	0.69 <sup>A</sup>
Age at diagnosis, mean (SD), years	31.7 (10.8)	37.3 (15.6)	31.6 (12.5)	34.5 (11.9)	0.77 <sup>B</sup>
Age at implant, mean (SD), years	33.0 (12.2)	38.6 (17.3)	35.5 (12.8)	37.5 (12.8)	0.83 <sup>B</sup>
Implant LVEF, mean (SD), %	25.6 (13.3)	19.7 (9.5)	20.8 (10.1)	18.8 (9.6)	0.58 <sup>B</sup>
Implant FS, mean (SD), %	10.3 (4.6)	8.5 (3.4)	9.3 (3.9)	8.6 (4.2)	0.90 <sup>B</sup>
Implant LVEDD, mean (SD), mm	69.8 (6.6)	72.3 (10.4)	73.1 (14.3)	71.7 (9.6)	0.95 <sup>B</sup>
Time on LVAD, mean (SD), days	214 (125)	212 (313)	317 (151)	520 (532)	0.11 <sup>B</sup>
Explant LVEF, mean (SD), %	64.0 (4.2)	n/a	65.9 (9.5)	n/a	0.64 <sup>C</sup>
Explant FS, mean (SD), %	29.5 (3.3)	n/a	31.9 (7.3)	n/a	0.53 <sup>C</sup>
Explant LVEDD, mean (SD), mm	44.5 (6.4)	n/a	54.3 (8.9)	n/a	0.05 <sup>C</sup>
<p><b>TTNtvs in cohort who recovered:</b> c.87624C&gt;A*; c.49346-1G&gt;A*; c.76383_76386delTAAT*; c.46782C&gt;A*; c.81518delC*; c.71326G&gt;T.</p> <p><b>TTNtvs in cohort who did not recover:</b> c.69976G&gt;T; c.67495C&gt;T*; c.41641C&gt;T; c.58172delA*.</p> <p>Titin variant position is given according to locus reference genomic (LRG) sequence 391_t1. A detailed overview of TTN gene structure, including the isoforms and protein domains affected by the TTNtvs described here, can be found at <a href="http://cardiodb.org/titin">http://cardiodb.org/titin</a>. *Variants reported in Roberts <i>et al</i><sup>2</sup>. P-values calculated with Fisher's exact test<sup>A</sup>, ANOVA<sup>B</sup> and unpaired t-test<sup>C</sup>.</p> <p><b>Abbreviations:</b> TTNtv=titin truncating variant, LVAD=left ventricular assist device, DCM=dilated cardiomyopathy, PPCM=peri-partum cardiomyopathy, LVEF=left ventricular ejection fraction, FS=fractional shortening, LVEDD=left ventricular end diastolic dimension, n/a=not applicable.</p>					



**Figure 1.**

