Supplementary Index

Use of Donor-Derived-Cell-Free DNA as a Marker of Early Allograft Injury in Primary Graft Dysfunction (PGD) to Predict the Risk of Chronic Lung Allograft Dysfunction (CLAD)

Michael Keller, Errol Bush, Joshua M. Diamond, Pali Shah, Joby Matthews, Anne W Brown, Junfeng Sun, Irina Timofte, Hyesik Kong, Ilker Tunc, Helen Luikart, Aldo Iacono, Steven D. Nathan, Kiran K. Khush, Jonathan Orens, Moon Jang, Sean Agbor-Enoh



Supplementary Figure 1: Study Design. 99 patients had Day 3 %ddcfDNA results. 13 patients did not have available Day 3 CXR or ABG data and were excluded from the final analysis. 86 patients were included in the final analysis.



Supplementary Figure 2: Differences in Day 3 %ddcfDNA between severity grades of PGD. Results demonstrate median, interquartile range, maximum and minimum values. Non PGD vs PGD 1+2 vs PGD 3 (8.5% (5.6, 13.2) vs 10.9% (8.20, 19.3) vs 17.6% (8.1, 22.8), p = 0.03)

Supplementary Table 1: Characteristics of Patients with CLAD vs without CLAD			
	No CLAD (N=56)	CLAD (N=22)	P Value
Sex at Birth: Male Female	23 (50%) 23 (50%)	8 (36%) 14 (66%)	0.43
Race: White Non-White	52 (93%) 4 (7%)	15 (68%) 7 (32%)	0.08
Type of Transplant: Single Double	18 (32%) 38 (68%)	5 (23%) 17 (77%)	0.45
Donor-Recipient Race Mismatch: Yes No	15 (27%) 41 (73%)	5 (33%) 15 (66%)	0.85
PGD Yes No	30 (49%) 26 (51%)	11 (50%) 11 (50%)	0.78
PGD 3 Yes No	9 (16%) 47 (84%)	3 (14%) 19 (86%)	0.79

Supplementary Table 1: Characteristics of patients with CLAD and without CLAD with corresponding univariate analysis. Only Race was incorporated into the multivariate analysis.

Race and PGD

Race+ PGD+ log2(%ddcfDNA) + PGD*log2(%ddcfDNA



Supplementary Figure 3a

Supplementary Figure 3b



Race+ PGD+ ACR + Infection + log2(%ddcfDNA) + PGD*log2(%ddcfDNA

Supplementary Figure 3c

Supplementary Figure 3. (a). ROC curve of multivariable model incorporating Race and PGD as predictors of CLAD. (b). ROC curve of multivariable model incorporating Race and PGD in addition to %ddcfDNA. (c). ROC curve of multivariate model incorporating Race, PGD, %ddcfDNA, Infection and Rejection.