

Table S1. Exome sequencing statistics.

	DLI	FL donor	FL recipient
Uniquely Mapping Statistics			
Mapreads	42406242	43886630	28608952
Bfast	30105898	30682229	22714177
Coverage Statistics			
		mapreads	
Total Bases Covered	33604042	33335470	32435260
-Average Depth	34.23	34.76	33.2
-Median Depth	18	17	12
-% Depth >= 10x	67.22%	66.10%	56.11%
Variant Statistics			
		mapreads	
Total SNVs	56139	52349	46043
-dbSNP	51187	47851	41990
-Missense	6859	6747	6286
-Stop Gain	57	62	54
-Stop Loss	10	9	12
-Start Loss	10	10	8
-Exon Junction	98	93	94
InDel Statistics			
		bfast	
Total Indels	42018	33749	63401
-Post Filtering*	5813	3794	4776
-dbSNP	202	148	138

*coding sequence (CDS), coverage >5 reads, SNP quality >5, consensus quality >5, mapping quality > 15, percent evidence > 15%