

Table S3A. Summary of Sanger sequencing validation of selected gene mutations (SNVs).

SNVs	FL donor			FL recipient		
	total no. selected variant calls	no. of validated variants	no. of somatic variants	total no. selected variant calls	no. of validated variants	no. of somatic variants
tier1	16	15	9	16	15	9
tier2	6	4	0	9	3	0
tier3	5	3	3	7	6	6
tier4	19	0	0	18	3	1
total	46	22	12	50	27	16

Table S3B. Summary of Sanger sequencing validation of selected gene mutations (InDels).

InDels	FL donor			FL recipient		
	total no. selected variant calls	no. of validated variants	no. of somatic variants	total no. selected variant calls	no. of validated variants	no. of somatic variants
tier1	14	4	2	14	4	2
tier2	11	0	0	9	0	0
tier3	7	1	0	5	1	0
total	32	5	2	28	5	2

tier1 identical variants in both lymphomas

tier2 different variants in identical genes in both lymphomas

tier3 variants unique to one lymphoma, high ranked QC

tier4 variants unique to one lymphoma, intermediate ranked QC