

# Table S2

## Oncogene exome analysis of iPSC vs donor PBMCs

- 1590\_101, no cancerous mutations in clone
- 1590\_103, no cancerous mutations in clone
- 1590\_104, no cancerous mutations in clone; seven sequence changes in clone
- 1592\_101, no cancerous mutations in clone; one heterozygous splicing change in DNTM3A. This variant is present at a frequency of 0.012 (43 / 3743) in the PBMC donor.
- 1592\_122, no cancerous mutations in clone
- 1592\_123, no cancerous mutations in clone
- 1620\_102, no cancerous mutations in clone; one variant with a one base pair deletion with 0.0261 frequency in the clone and 0.0268 frequency in the donor PBMC
- 1620\_104, no cancerous mutations in clone

TUTE predicted deleterious mutations novel in the clones

### *1590\_101, no cancerous mutations in clone*

Code

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Search:

CHROM	POS	REF	ALT	GT.x	GT.y	AF.x	AF.y	DP.x	INFO.x
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### *1590\_103, no cancerous mutations in clone*

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### *1590\_104, no cancerous mutations in clone; seven new changes in clone*

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