The following are the setting used process raw data into the plots shown in the manuscript.

**Figure 2: Genomic instability test**

The file: stabilty\_example.csv

Model: Instability

Targets for normalization ( chromosome (first ‘delta’) ): CHR4

Target order: CHR1,CHR4,CHR8,CHR10,CHR12,CHR17,CHR18,CHR20,CHRX

Sample order: GM25953,GM25975,GM25974,GM25952,Normal

Control sample Second ‘delta’: Normal

**Figure 3: Absolute quantification**

The file: 5 files in the folder ‘Absolute

-- All files can be uploaded at once

-- The reference gene must be included

Model: Absolute

Targets for normalization (endogenous control genes): ACTB,GAPDH

Target order: left blank

*Control cell lines and stages order ------ []*

Sample order: NCRM1-IPSC,522-266-2-IPSC,AiW001-2-IPSC,AiW002-2-IPSC,AJC001-5-IPSC,AJG001C4-IPSC,NCRM1-NPC,522-266-2-NPC,AiW001-2-NPC,AiW002-2-NPC,AJC001-5-NPC, AJG001C4-NPC,NCRM1-DA4W,522-266-2-DA4W,AiW001-2-DA4W,AiW002-2-DA4W,AJG001C4-DA4W,AJC001-5-DA4W,NCRM1-DA6W,522-266-2-DA6W,AiW001-2-DA6W,AiW002-2-DA6W,AJG001C4-DA6W,AJC001-5-DA6W

*Control order of stages ----- [works]*

Sample order : IPSC,NPC,DA4W,DA6W

*Control cell lines order ----- [works]*

Sample order: NCRM1,522-266-2,AiW001-2,AiW002-2,AJC001-5,AJG001C4

statistic –

number of groups: 4

Groups are defined within the sample name column.

order :

IPSC,NPC,DA4W,DA6W

Selection for time series or repeated measures/ linked samples : yes

Selection for parametric or normal distribution: yes

**Figure 4: Relative quantification be two method**

**Relative quantification delta CT model:**

The file: RELATIVE\_example.csv

Model: Relative (delta CT)

Targets for normalization (endogenous control genes): ACTB,GAPDH

Targets order (genes): PAX6,CAMK2A,GRIN1

Sampel order (cell lines, time points): AiW002-2-D0, AiW002-2-D7,KYOU-D0,KYOU-D7