

## CORRIGENDUM

## Sporadic and reversible chromothripsis in chronic lymphocytic leukemia revealed by longitudinal genomic analysis

L Bassaganyas, S Beà, G Escaramís, C Tornador, I Salaverria, L Zapata, O Drechsel, PG Ferreira, B Rodriguez-Santiago, JMC Tubio, A Navarro, D Martín-García, C López, A Martínez-Trillos, A López-Guillermo, M Gut, S Ossowski, C López-Otín, E Campo and X Estivill

*Leukemia* (2015) 29, 758; doi:10.1038/leu.2014.309**Correction to:** *Leukemia* (2013) 27, 2376–2379; doi:10.1038/leu.2013.127

Since the publication of this article, the authors have identified several errors within their paper. These are listed as follows:

Table 1:

Column 'AAChange':

- First row (without counting Header) should be NM\_003902:c.C1558T:p.Q541X
- Second row should be NM\_001127710:c.A2299G:p.T901A
- Third row should be NM\_001080412:c.C1924T:p.P643S

Supplementary Table S1:

In the 'ID sample' column, 'Sample Year' 2006 is correct but ID sample should be 016-T06.

Supplementary Table S3:

- Row 10 (without counting Header) column 'locus' should be t(1;10)(q32;q24).
- Row 24 (without counting Header) column 'locus' should be t(6;10)(q15;24).
- Row 50 (without counting Header) column 'brk 2' should be 19284700 and column 'size' should be 19166140.

The corrected Table 1 is displayed here, and the corrected Supplementary Tables accompany the online version of this corrigendum.

The authors would like to apologise for any inconvenience this may have caused.

**Table 1.** Somatic point mutations in all time-point samples

| Chromosome | Position  | Reference | Observed | Gene     | Exonic function   | AAChange                      | 016-T00 | 016-T02 | 016-T05 | 016-T11 |
|------------|-----------|-----------|----------|----------|-------------------|-------------------------------|---------|---------|---------|---------|
| 1          | 78425887  | G         | A        | FUBP1    | stopgain SNV      | NM_003902:c.C1558T:p.Q541X    | +       | +       | +       | +       |
| 1          | 186277552 | A         | G        | PRG4     | nonsynonymous SNV | NM_001127710:c.A2299G:p.T901A | +       | +       | +       | +       |
| 3          | 141163154 | C         | T        | ZBTB38   | nonsynonymous SNV | NM_001080412:c.C1924T:p.P643S | +       | +       | +       | +       |
| 6          | 44230329  | C         | A        | NFKBIE   | stopgain SNV      | NM_004556:c.G853T:p.E285X     | –       | +       | –       | –       |
| 6          | 132892234 | A         | C        | TAAR6    | nonsynonymous SNV | NM_175067:c.A774C:p.K258N     | +       | +       | +       | +       |
| 6          | 146264389 | T         | C        | SHPRH    | nonsynonymous SNV | NM_001042683:c.A2128G:p.M710V | +       | +       | +       | +       |
| 11         | 108114749 | G         | C        | ATM      | nonsynonymous SNV | NM_000051:c.G566C:p.R189T     | +       | +       | +       | +       |
| 12         | 21358921  | C         | A        | SLCO1B1  | nonsynonymous SNV | NM_006446:c.C1451A:p.P484H    | +       | +       | +       | +       |
| 15         | 45409873  | A         | G        | DUOXA1   | nonsynonymous SNV | NM_144565:c.T1292C:p.V431A    | +       | +       | +       | +       |
| 16         | 745751    | T         | C        | FBXL16   | nonsynonymous SNV | NM_153350:c.A806G:p.N269S     | +       | +       | +       | +       |
| 17         | 42463002  | G         | A        | ITGA2B   | nonsynonymous SNV | NM_000419:c.C491T:p.A164V     | +       | +       | +       | +       |
| 19         | 19431945  | G         | A        | MAU2     | splicing site     | NA                            | +       | +       | +       | +       |
| 19         | 56173950  | A         | T        | U2AF2    | nonsynonymous SNV | NM_001012478:c.A569T:p.Q190L  | +       | +       | –       | +       |
| X          | 125299321 | C         | A        | DCAF12L2 | nonsynonymous SNV | NM_001013628:c.G587T:p.W196L  | +       | +       | +       | +       |

+, present; –, absent.