**Table File General Guidelines**

* Tables can also be included in the manuscript file itself or it can be attached separately as Table File in word format using .doc file or docx file.
* All tables should have a concise title and written as Table 1 with a period (.). E.g., Table 1. Stimulation settings
* Footnotes can be used to explain abbreviations.
* Tables extending beyond 1 page should be avoided.

Refer the Table file template given in the next page:

**Table File**

**Table 1.** *α*-Glucosidase inhibition of *Walsura trifoliata* hexane, ethyl acetate, methanol root extracts and acarbose

|  |  |  |  |
| --- | --- | --- | --- |
| **Sample** | **Concentration (µg/ml)** | **% of Inhibition** | **IC50(µg/ml)** |
| Hexane extract | 200 | 7.00±1.80 |  |
|  | 400 | 11.62±2.07 |  |
|  | 600 | 20.00±2.71 | 50 |
|  | 800 | 27.52±2.91 |  |
|  | 1000 | 31.45±1.56 |  |
| Ethyl acetate extract | 200 | 14.70±1.56 |  |
|  | 400 | 25.47±1.94 |  |
|  | 600 | 32.13±2.31 | 50 |
|  | 800 | 36.23±1.29 |  |
|  | 1000 | 43.41±2.07 |  |
| Methanol extract | 200 | 24.61±2.56 |  |
|  | 400 | 35.89±2.35 |  |
|  | 600 | 44.10±1.53 | 690.10±1.44 |
|  | 800 | 56.92±1.35 |  |
|  | 1000 | 65.47±2.31 |  |
| Acarbose | 200 | 46.49±1.29 |  |
|  | 400 | 54.87±2.35 |  |
|  | 600 | 65.98±2.52 | 290.90±1.82 |
|  | 800 | 72.82±2.56 |  |
|  | 1000 | 81.19±2.58 |  |

Each value represents the mean ± SEM of triplicate experiments.

Table 2a, b. Summary of chromosome region, on which subtelomere FISH probe localized, in leukemias and lymphomas with 11q22-q25 or 17p13 abnormality: (a)Telomere region at 11q22-q25 (subtelomere probe: VIJ22072)

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Type of chromosome | No. patients analyzed | Probe signal located on | | | Loss |
| Terminal | Central | Partner chromosome |
| t(11;α)(q22-q25;α) | 0(11) | ND | ND | ND | ND |
| add(11)(q22-q25) | 9(11) | 1 | 3 | 2 | 3 |
| del(11)(q22-q25) | 7(9) | 4 | 0 | 0 | 3 |
| r(11)(pter;qter) | 1(1) | 0 | 1\* | 0 | 1\* |
| Total No. analyzed | 17 (32)\*\* | 5 | 4 | 2 | 7(40.7%) |
| *\*The patient had two cell clones, in which one had signal located on ring chromosome and others had loss of signal. \*\* Numbers in parenthesis are total of observed patients. 15 out of 32 patients were not analyzed by subtelomere probe. ND: not observed, α: any partner chromosome* | | | | | |

(b)Telomere region at 17p13（subtelomere probe: D17S34）

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Type of chromosome | No. patients analyzed |  | | | Loss |
| Terminal central | Partner | Chromosome |
| t(17;α)(p13;α) | 6(6) | 0 | 0 | 4 | 2 |
| add(17)(p13) | 10(10) | 4 | 0 | 0 | 6 |
| del(17)(p13) | 8(8) | 0 | 0 | 0 | 8 |
| Total No. analyzed | 24(24)\*\* | 4 | 0 | 4 | 16(66.0%) |
| *\*\* Numbers in parenthesis are total of observed patients. α: any partner chromosome* | | | | | |