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|  | **Reviewer 1** | **Reviewer 2** | **Changes** |
| Findings of this study | Confirm known results | Make a significant contribution to existing knowledge. |  |
| Title of the abstract | Is appropriate to the thematic area and descriptive. | Is appropriate to the thematic area and descriptive. | Screening the Molecular Diversity of Selected Interim Clones of Rubber (*Hevea brasiliensis*) Using SSR Molecular Markers |
| Content of the abstract | Needs improvements.  Some sentences are needed to be restructuring as their meanings are not clear. | Is clear and concise. | Rearranged sentences   1. Among those, HB6, HB28, HB29, hmac4, and hmct1 SSR primers produced clear and detectable bands while HB4 and HB8 primers failed to produce clearly detectable bands. 2. This molecular analysis revealed that both Wickham and non-Wickham genetic bases that exists in the Sri Lankan breeding pool showed narrow genetic diversity |
| Recommendation | Accept with minor corrections.  1) Wickham and non-Wickham genetic bases  2) restructure this sentence "It extensively used the Wickham genetic base for developing the current breeding pool, it is believed that the current genetic diversity of rubber in Sri Lanka, is significantly narrowed hence;" | Accept with minor corrections.  The abstract can be accepted with some (minor) language editing. | 2) Most of the parental lines utilized in the ongoing breeding programs belong to Wickham genetic base. Because it is believed that the current genetic diversity of rubber in Sri Lanka, is considerably narrow and it has reached to its threshold level for economically important characteristics. |