

ABSTRACT

Disclosed are a sequence report method and a sequence report device for reducing a signaling amount for reporting a Zadoff-Chu sequence or a GCL sequence allocated for a cell. Indexes starting at 1 are correlated to different ZC sequences and are allocated for cells so that the indexes are continuous. When such ZC sequences are reported from BS to UE, a start index indicating the start of the continuous indexes is combined with the number of allocated sequences and they are reported as allocation sequence information by a report channel. The UE and the BS share the correlation between the ZC sequences and the indexes and the UE identifies a usable sequence number according to the correlation and the allocation sequence information reported from the BS.