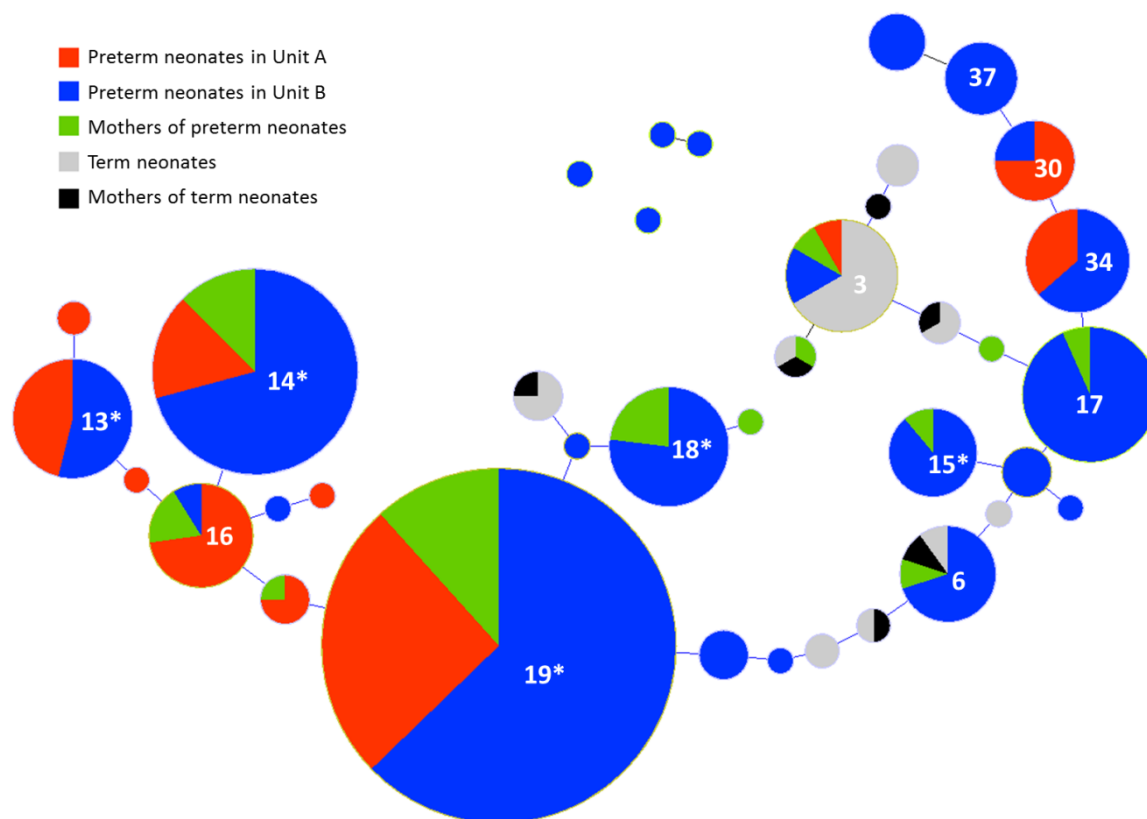


**Supplemental Digital Content 3.** goeBURST analysis of 37 distinct MLVA-types of 227 *S. haemolyticus* isolates from skin and gut of neonates and breast milk of their mothers at single locus variant level.



MLVA-types (n=7) and isolates (n=16) yielding two bands in loci SH0999 or SH0324 were excluded. MLVA-types that colonized at least 5 neonates are depicted by their MLVA-type numbers. Node represents a distinct MLVA-type, whereas the size of the node is proportional to the number of isolates of that MLVA-type. Solid line indicates relatedness of the MLVA-types at single locus variant level. The color of the node indicates the source of isolation of the MLVA-type: red – preterm neonates in Unit A; blue – preterm neonates in Unit B; green – mothers of preterm neonates; grey – term neonates; black – mothers of term neonates. MLVA-types isolated from bloodstream of neonates with late-onset sepsis are denoted by white asterisk.