igem_part

no <- c(1, 2) team_name <- c("Queens-Canada", "Tainan") project_title <- c("Velcrion", "Oh_My_Gut") project_year <- c(2020, 2019) wiki_page <- c("https://2020.igem.org/Team:Queens_Canada", "https://2019.igem.org/Team:NCKU_Tainan")

igem_team <- data.frame(no, team_name, project_title, project_year, wiki_page) #knitr::kable(std, format = "markdown") knitr::kable(igem_team)

no <- c(1, 2, 3)

part_id <- c("BBa_K2259000", "BBa_K2259010", "BBa_K2259092") part_name <- c("SynORI framework RNA II - Replication Initiator (Group A)", "Rop protein - global copy number inhibitor (SynORI framework)", "Minimal base vector for SynORI system building") part_type <- c("Prject", "Coding", "Plasmid") team_id <- c(2, 2, 2)

igem_part <- data.frame(no, part_id, part_name, part_type, team_id)

knitr::kable(igem_part)