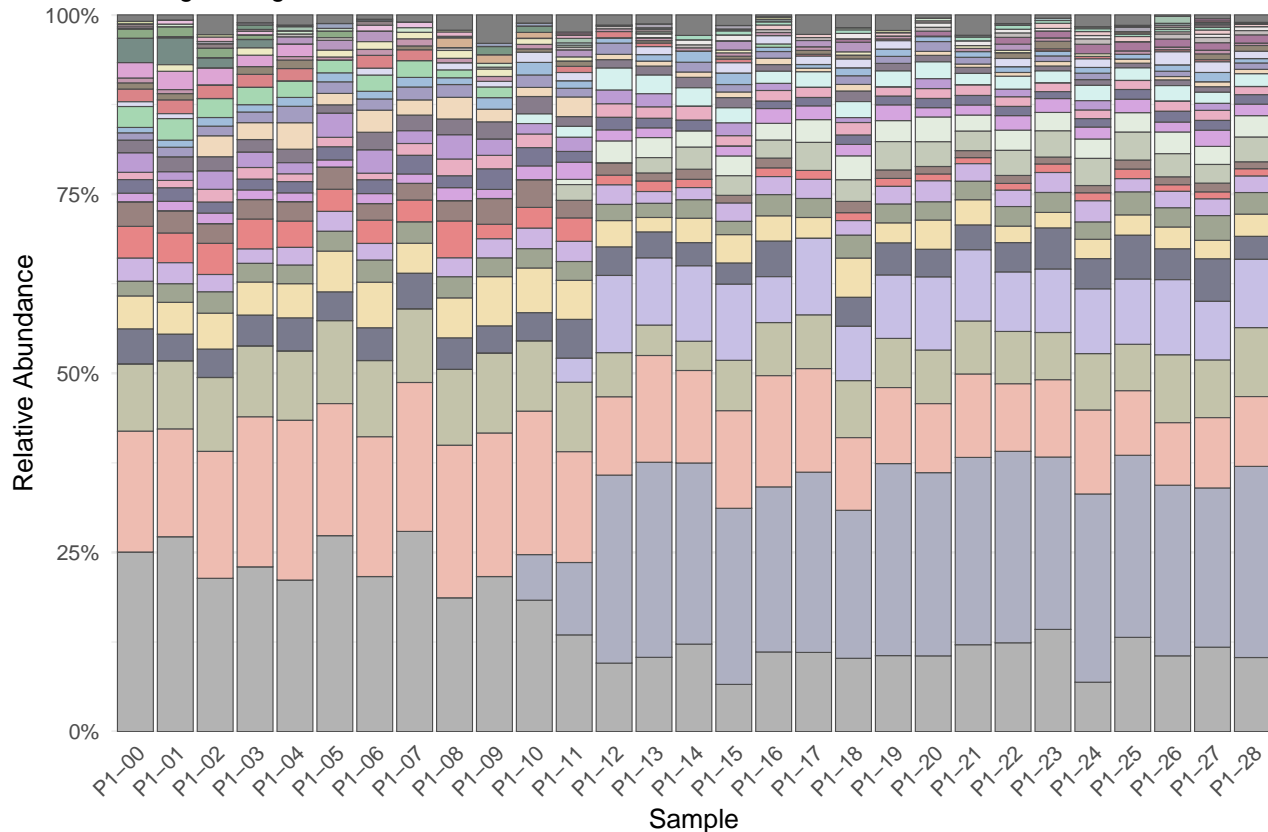


# Community composition – top 50 clusters

Average linkage at 3.0% → 180 clusters



## Cluster (top-hit species)

CL156 – JEMV_s	CL29 – Enterobacter kobei
CL109 – CP011270_s	CL30 – ALOU_s
CL82 – Pseudobacter ginsenosidimutans	CL26 – Pedobacter ginsenosidimutans
CL99 – Bacillus mesonae	CL24 – Terrabacter korensis
CL67 – Pseudoxanthomonas japonensis	CL20 – Lysobacter korensis
CL78 – Bosea vaviloviae	CL19 – Prosthecobacter fluviatilis
CL45 – Delftia acidovorans	CL16 – Flaviaesturariibacter terrae
CL66 – EU135464_s	CL17 – Herminiimonas contaminans
CL58 – Delftia acidovorans	CL21 – Ensifer adhaerens
CL57 – Lysobacter panacisoli	CL15 – CP012155_s
CL41 – Chitinophaga eiseniae	CL18 – Acinetobacter oleivorans
CL36 – Enterobacter kobei	CL14 – Flavobacterium lindsaytolerans
CL60 – Pseudobacter ginsenosidimutans	CL12 – Herbaspirillum seropedicae
CL53 – Pseudoxanthomonas indica	CL8 – Novosphingobium endophyticum
CL35 – HQ119897_s	CL13 – Chryseobacterium cucumeris
CL31 – JQJY_s	CL10 – Sphingomonas kyeonggiensis
CL25 – Paraburkholderia graminis	CL11 – Pseudomonas granadensis
CL43 – Bosea vaviloviae	CL7 – Flavobacterium brevitiae
CL28 – Labrys neptuniae	CL5 – Acidovorax delafeldii
CL27 – CP003911_s	CL6 – Pedobacter zeaxanthinifaciens
CL22 – Flavobacterium defluvi	CL9 – Cupriavidus plantarum
CL33 – Pseudoduganella eburnea	CL3 – Pseudomonas frederiksbergensis
CL23 – Duganella ginsengisoli	CL4 – Paenarthrobacter nitroguajacolicus
CL42 – Dyadobacter fermentans	CL2 – Sphingobium czechense
CL32 – PEBF_s	CL1 – Bacillus cucumis

Clusters not in top 50 are grouped as "Other".