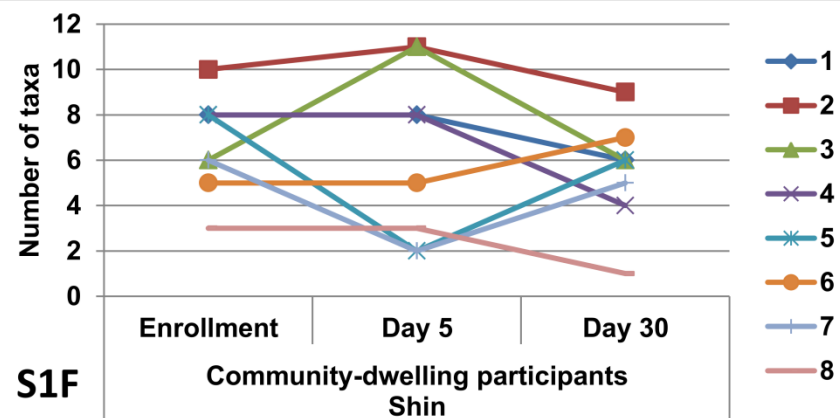
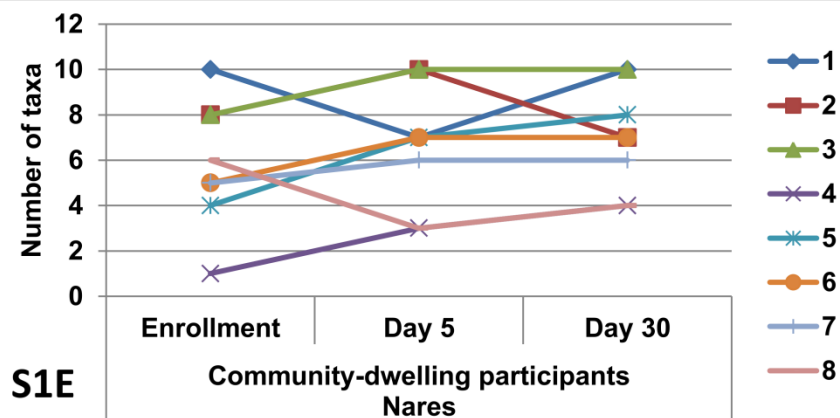
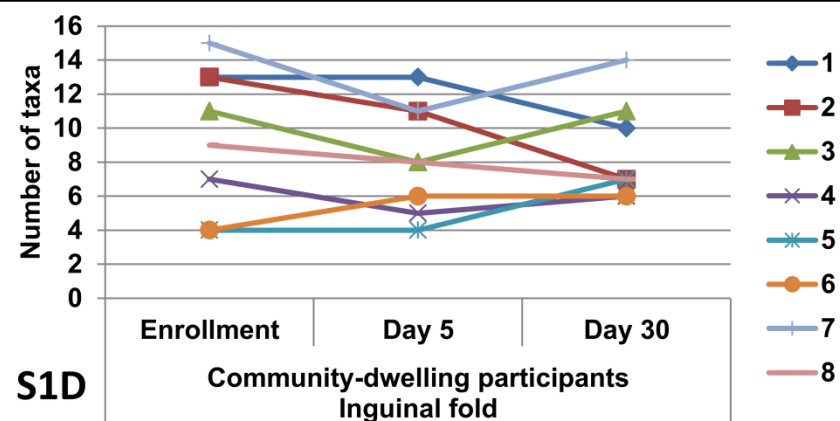
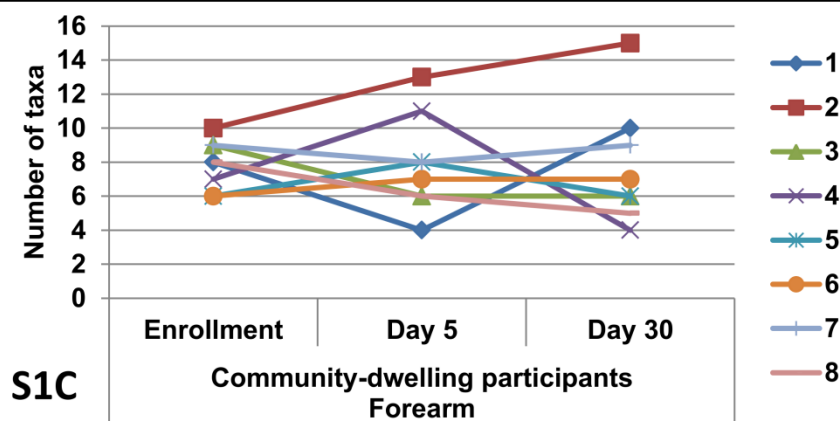
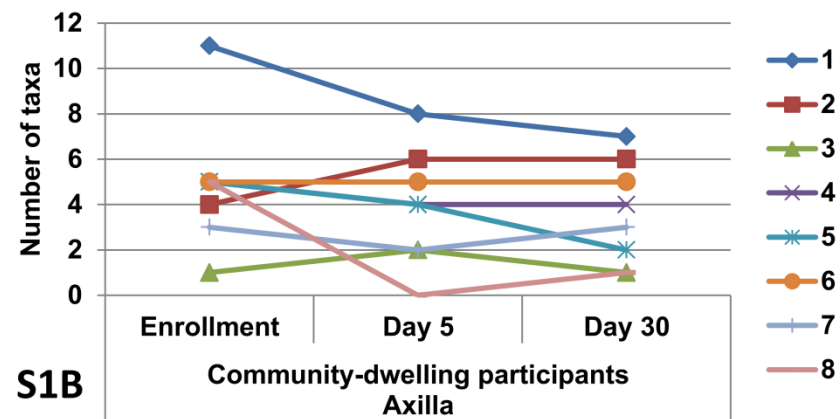
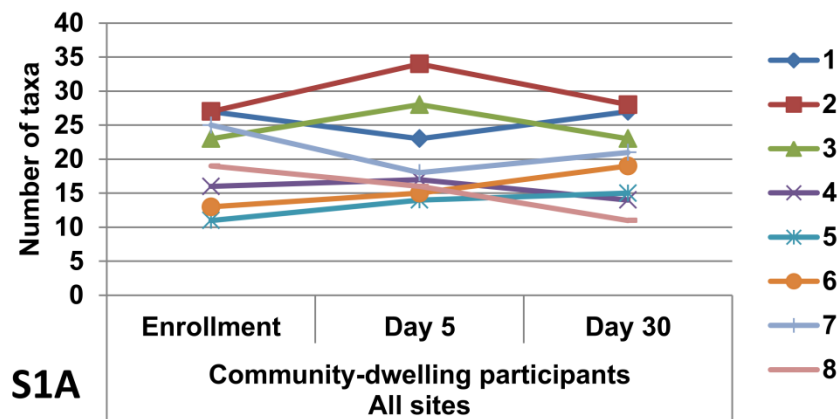


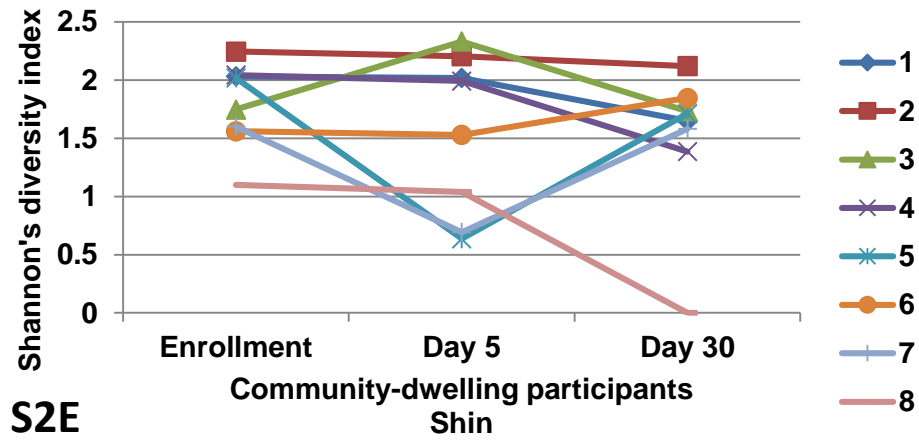
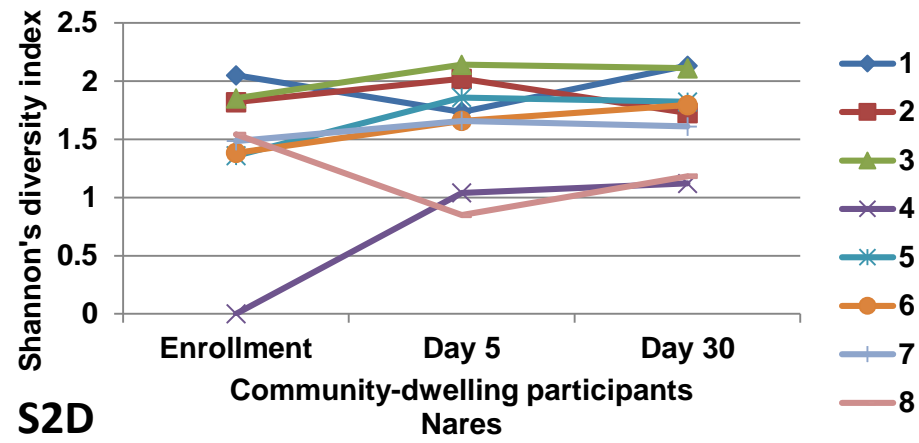
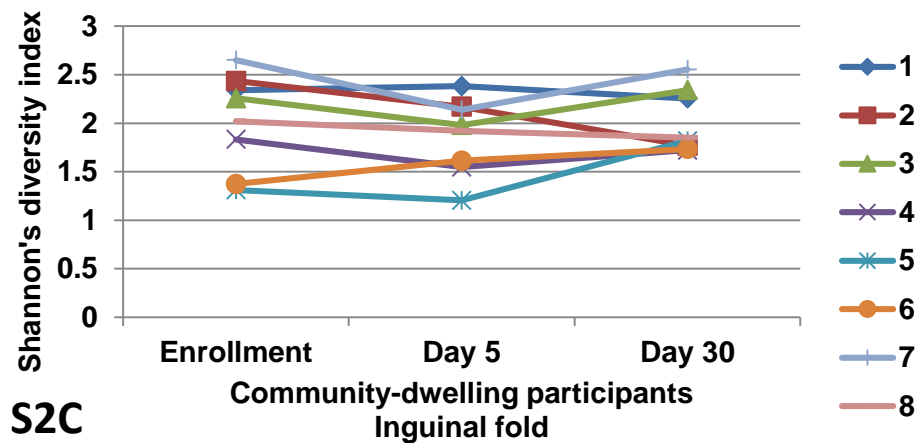
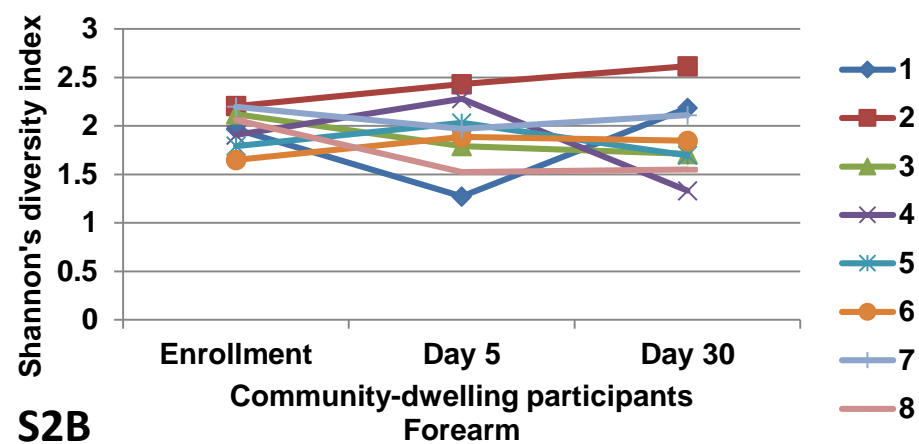
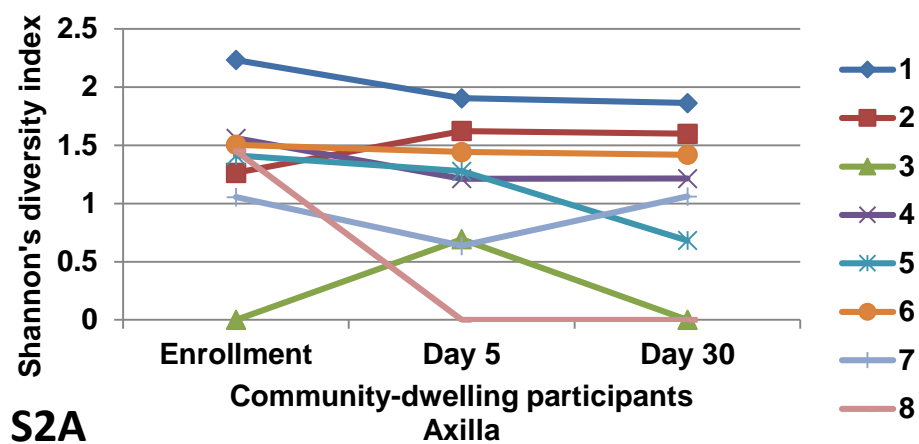
**Table S1. Culture media and incubation environments.**

<b>Media<sup>a</sup></b>	<b>Incubation Conditions</b>	<b>Incubation Interval</b>	<b>Volume Plated</b>
Trypticase soy agar with 5% sheep blood	35 °C, 5% CO <sub>2</sub>	Three days (first read after 24 hours)	10 uL and 100 uL
Chocolate agar	35 °C, 5% CO <sub>2</sub>	Three days (first read after 24 hours)	50 uL
Colistin naladixic acid agar	35 °C, 5% CO <sub>2</sub>	Three days (first read after 24 hours)	50 uL
MacConkey agar	35 °C, O <sub>2</sub>	Three days	50 uL
Chopped meat broth enrichment	35 °C, Anaerobic jar	Incubate for 24 hours, then subculture to BBA, BBE, LKV	50 uL
<i>Brucella</i> blood agar with hemin and vitamin K (BBA)	35 °C, Anaerobic jar	Seven days (first read after 48 hours)	100 uL and 10 uL
<i>Bacteroides</i> bile esculin (BBE)	35 °C, Anaerobic jar	Seven days (first read after 48 hours)	50 uL
<i>Brucella</i> laked blood agar with kanamycin and vancomycin (LKV)	35 °C, Anaerobic jar	Seven days (first read after 48 hours)	50 uL
Sabbaroud's agar with chloramphenicol	25 °C, O <sub>2</sub>	Three weeks (first read after 48 hours, plates sealed after 48 hours to prevent dehydration)	50 uL

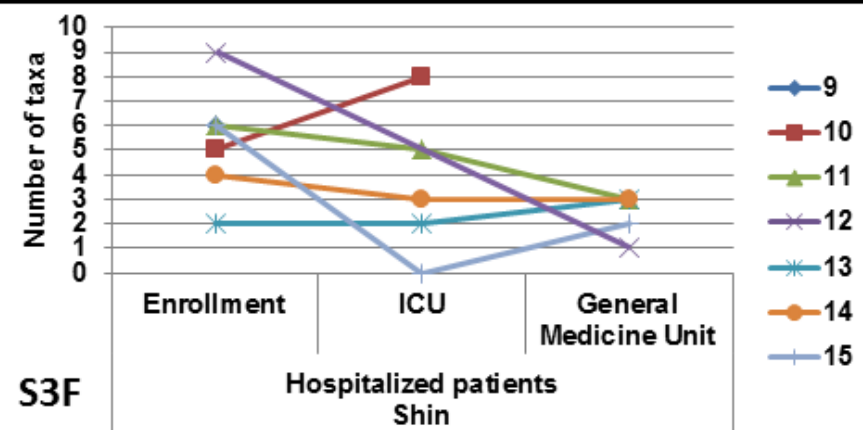
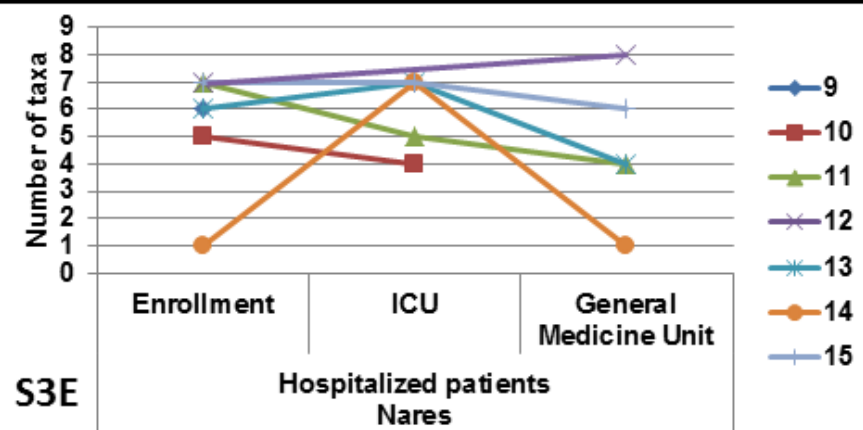
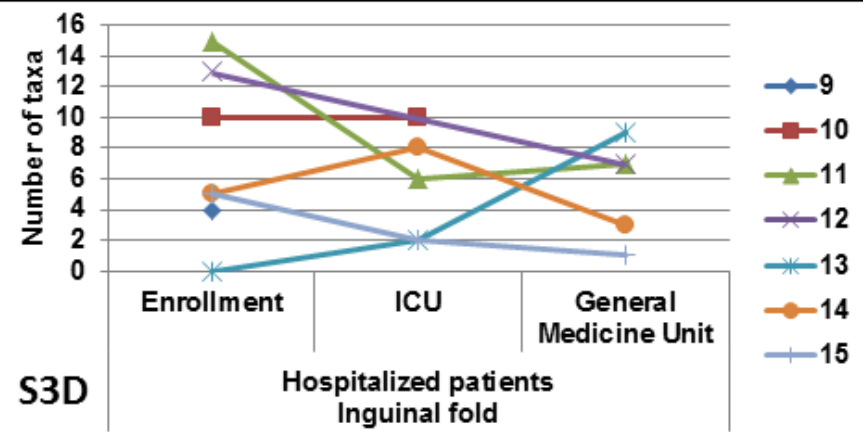
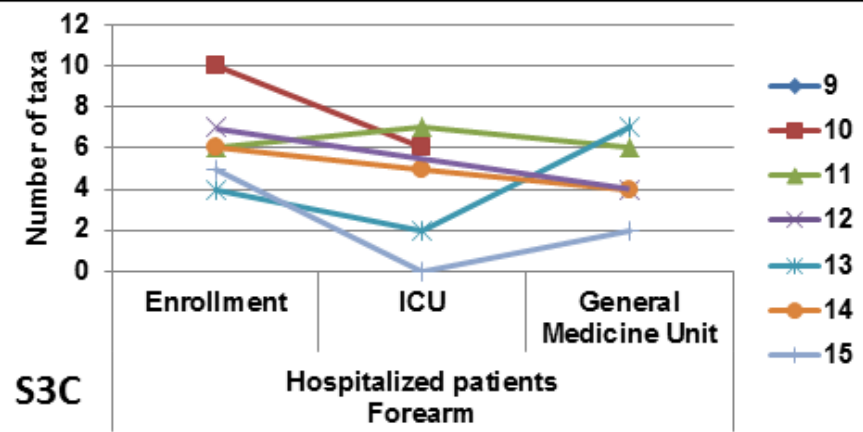
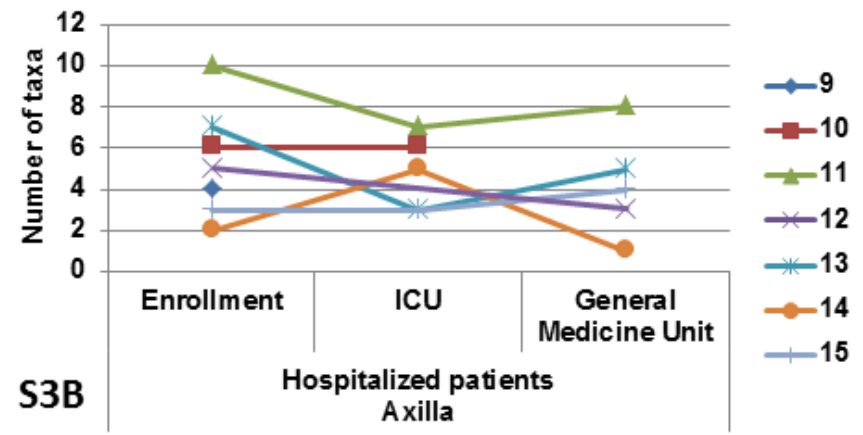
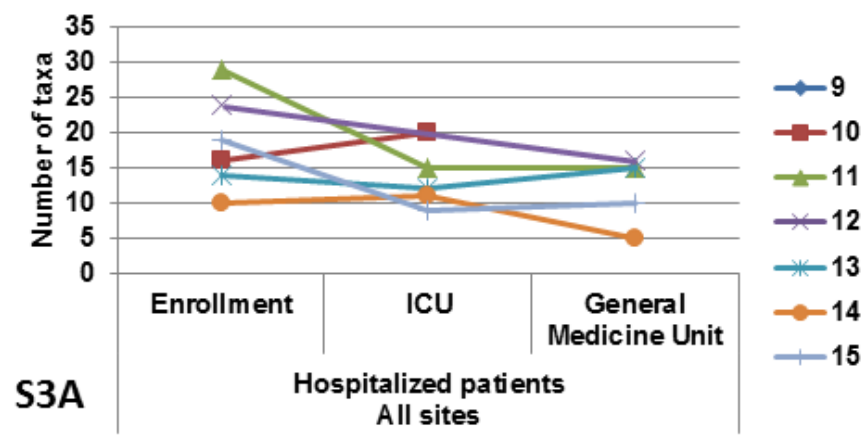
<sup>a</sup> All medium from Hardy Diagnostics



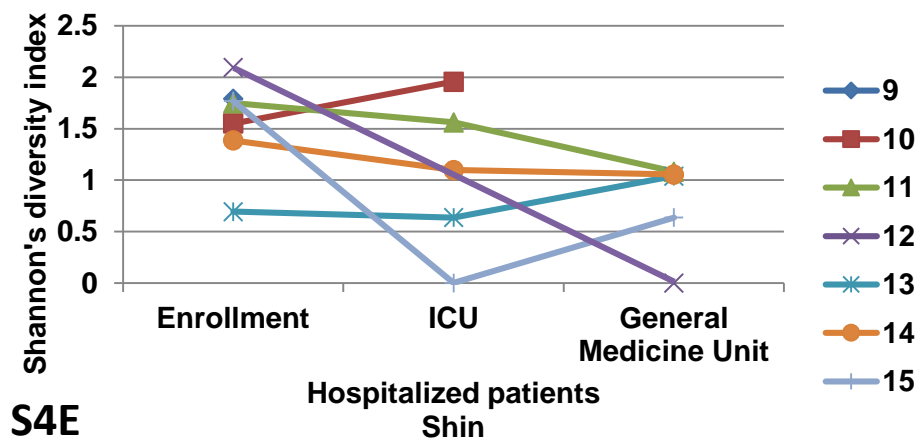
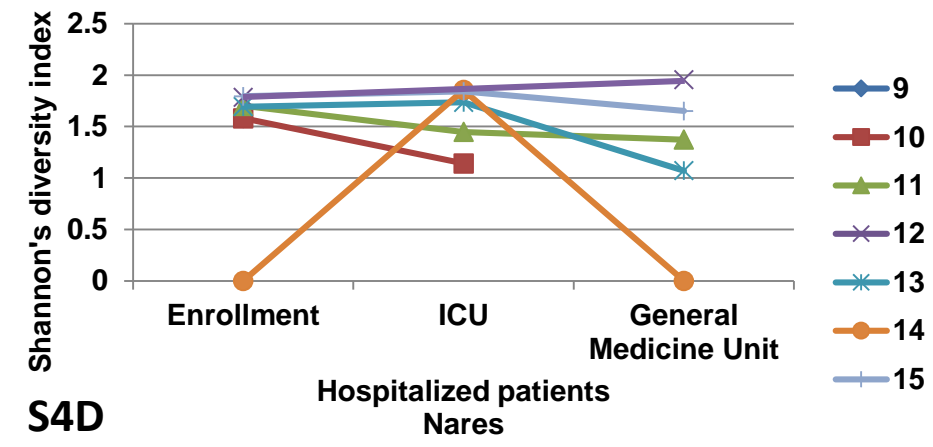
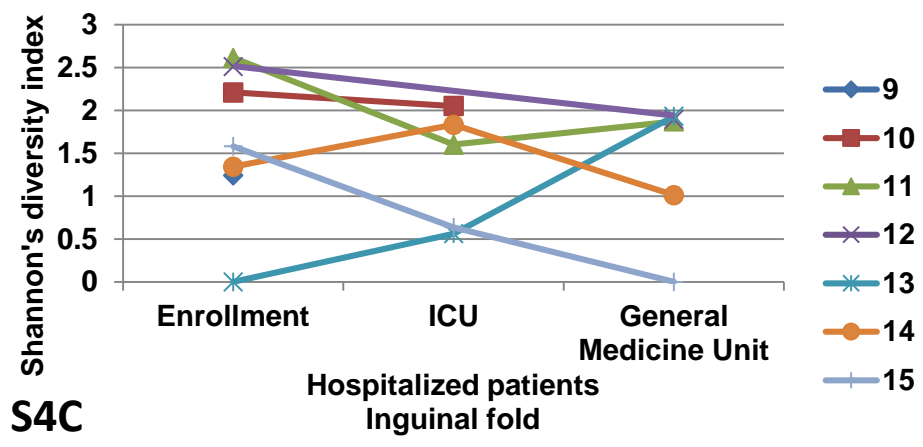
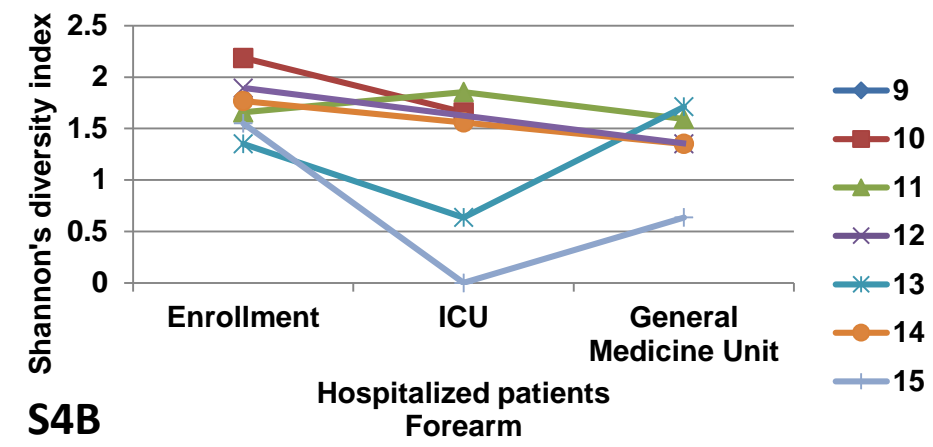
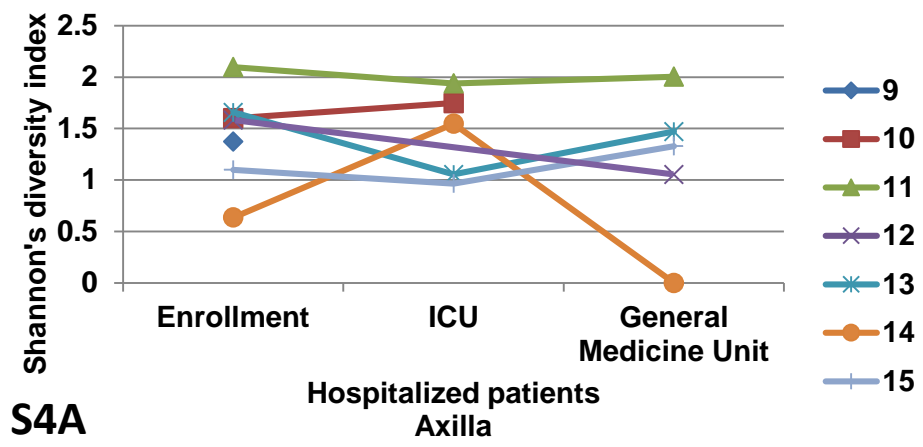
**Figure S1.** Richness of each community-dwelling participant at each time point. Each line represents a distinct participant. **A.** All sites combined. **B.** Axilla. **C.** Forearm. **D.** Inguinal fold. **E.** Nares. **F.** Shin.



**Figure S2.** Diversity of each community-dwelling participant at each time point. Each line represents a distinct participant. **A.** Axilla. **B.** Forearm. **C.** Inguinal fold. **D.** Nares. **E.** Shin.



**Figure S3.** Richness of each hospitalized patient at each time point. Each line represents a distinct patient. **A.** All sites combined. **B.** Axilla. **C.** Forearm. **D.** Inguinal fold. **E.** Nares. **F.** Shin.



**Figure S4.** Diversity of each hospitalized patient at each time point. Each line represents a distinct patient. **A.** Axilla. **B.** Forearm. **C.** Inguinal fold. **D.** Nares. **E.** Shin.