

Microbiome beta diversity, Bray-Curtis, genus Principal Coordinates Analysis plot 1.0 -XT_D10=NFlex=R1 XT_904=NFIex=R1=€ Flex D04-NF-R1 Flex D10-NF-R1 Flex D03-NF-R1 XT_B03=NF18X=R1=8 0.5 XT_D02-NFlex-R1-B Flex_D02-NF-R1 PC2 (18.9% of variance) Group Flex XT 0.0 PARTIES FRILES XT_D09-NFIe Flex_D09-NF-R1 XT_D06-NIEWDRA-MF-R1 Flex D06-NF-R1 -0.5FIEXT DOOF NITE BY-R1-B Flex TO ON THE REAL POINTS

0.0

PC1 (43.1% of variance)

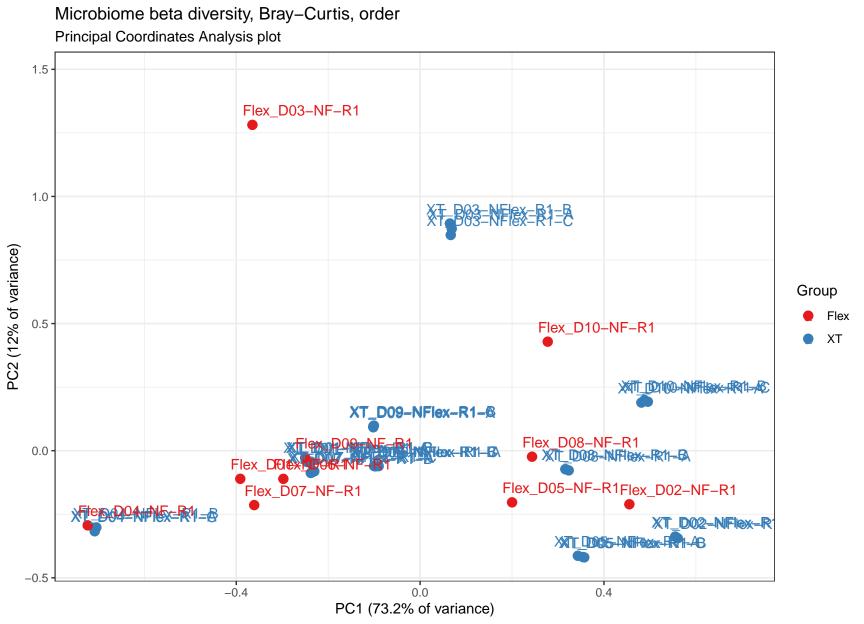
0.5

1.0

-0.5

-1.0

Microbiome beta diversity, Bray-Curtis, family Principal Coordinates Analysis plot 1.0 Flex_DOT-NFIEX-R1-8 XT_D07-NF16x-1R1-18 Flex_D07-NF-R1 0.5 Flex_D06-NF-R1 PC2 (25.4% of variance) Flex_D08-NF-R1 Flex_D09-NF-Group XT_B88=NFlex-R1=R Flex XT_D09-NEIEXOB3-NEIEXRATAB XT 0.0 Flex_D05-NF-R1 FlexXID020NFNRex-R1 Flex D04-NF-R1 Flex_D10-NF-R1 XT_D110=NF1ex=R -0.5 0.0 0.5 PC1 (48.2% of variance)



Microbiome beta diversity, Bray-Curtis, class Principal Coordinates Analysis plot 0.5 XT_D02-NFIex-R XT_1005-NF16xx-002-18F-R1 Flex D07-NF-R1 XT_D08-NFIbx-R1-B Flex_DOM Flow R11-AB Flex 06-NFT 1006-NFTex-PR11-0.0 Flex_D09-NF-R1 XT_D09-NFlex-R1-8 SA-TIPAL WHITEHAM - CONTINUE TO A CONTINUE T PC2 (11.8% of variance) Group Flex_D10-NF-R1 Flex XT -0.5 -1.0Flex_D03-NF-R1 -0.40.0 0.4 PC1 (73.6% of variance)

