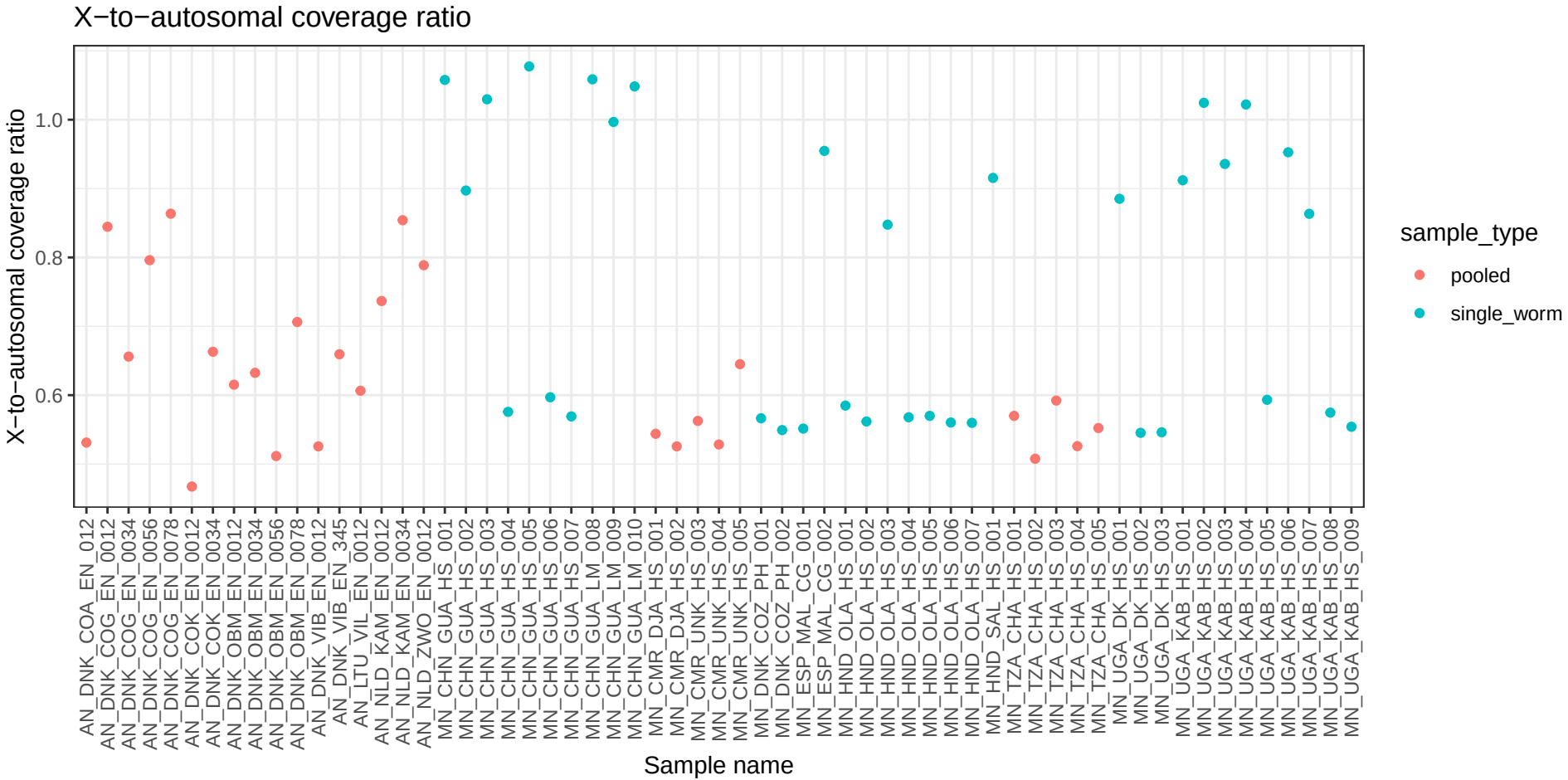
X−to−autosomal coverage ratio
X−to−autosomal coverage ratio
1.0
0.8
0.6
AN_DNK_COA_EN_012
AN_DNK_COG_EN_0012
AN_DNK_COG_EN_0034
AN_DNK_COG_EN_0056
AN_DNK_COG_EN_0078
AN_DNK_COK_EN_0012
AN_DNK_COK_EN_0034
AN_DNK_OBM_EN_0012
AN_DNK_OBM_EN_0034
AN_DNK_OBM_EN_0056
AN_DNK_OBM_EN_0078
AN_DNK_VIB_EN_0012
AN_DNK_VIB_EN_345
AN_LTU_VIL_EN_0012
AN_NLD_KAM_EN_0012
AN_NLD_KAM_EN_0034
AN_NLD_ZWO_EN_0012
MN_CHN_GUA_HS_001
MN_CHN_GUA_HS_002
MN_CHN_GUA_HS_003
MN_CHN_GUA_HS_004
MN_CHN_GUA_HS_005
MN_CHN_GUA_HS_006
MN_CHN_GUA_HS_007
MN_CHN_GUA_LM_008
MN_CHN_GUA_LM_009
MN_CHN_GUA_LM_010
MN_CMR_DJA_HS_001
MN_CMR_DJA_HS_002
MN_CMR_UNK_HS_003
MN_CMR_UNK_HS_004
MN_CMR_UNK_HS_005
MN_DNK_COZ_PH_001
MN_DNK_COZ_PH_002
MN_ESP_MAL_CG_001
MN_ESP_MAL_CG_002
MN_HND_OLA_HS_001
MN_HND_OLA_HS_002
MN_HND_OLA_HS_003
MN_HND_OLA_HS_004
MN_HND_OLA_HS_005
MN_HND_OLA_HS_006
MN_HND_OLA_HS_007
MN_HND_SAL_HS_001
MN_TZA_CHA_HS_001
MN_TZA_CHA_HS_002
MN_TZA_CHA_HS_003
MN_TZA_CHA_HS_004
MN_TZA_CHA_HS_005
MN_UGA_DK_HS_001
MN_UGA_DK_HS_002
MN_UGA_DK_HS_003
MN_UGA_KAB_HS_001
MN_UGA_KAB_HS_002
MN_UGA_KAB_HS_003
MN_UGA_KAB_HS_004
MN_UGA_KAB_HS_005
MN_UGA_KAB_HS_006
MN_UGA_KAB_HS_007
MN_UGA_KAB_HS_008
MN_UGA_KAB_HS_009
Sample name
sample_type
pooled
single_worm